SEARCH REQUEST FORM

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PTO-1590 (9-90)

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GenCore version 4.5

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OM protein - protein search, using sw model

Run on: May 22, 2002, 16:16:24; Search time 29:93 Seconds

(without alignments)
578.935 Million cell updates/sec

Perfect score: 156
Sequence: 1 MAYLSGALGFRMKDSALKVL......IPEDPAWDAPITDFYFQQCD 156
Scoring table: OLIGO
Gapop 60.0, Gapext 60.0
Searched: 747574 seqs, 111073796 residues

If size: 12
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Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Total number of hits satisfying chosen parameters:

39

Post-processing: Listing first 1000 summaries

Database :

A_Geneseq_032802:* : /SIDSI/gcdpats_hold-geneseg/genesegp-embl/AA1980.DAT:	/sibsi/ycydata/noid-geneseq/geneseqp-embl/AA1981.DAT: /SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1982.DAT	/SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1983.DAT:*	'SIDSI/gcgdata/hold-geneseq/geneseqp-embl/AA1984.DAT:*	'SIDSI/gcgdata/hold-geneseq/geneseqp-embl/AA1985.DAT:*	'SIDSI/gcgdata/hold-geneseq/geneseqp-embl/AA1986.DAT:*	'SIDSI/gcgdata/hold-geneseq/geneseqp-embl/AA1987.DAT:	/SIDSI/gcgdata/hold-geneseq/geneseqp-embl/AA1988.DAT:*	/SIDSI/gcgdata/hold-geneseq/geneseqp-embl/AA1989.DAT:*	/SIDS1/gcgdata/hold-geneseq/genesegp-emb1/AA1990.DAT:*	/SIDS1/gcgdata/hold-geneseq/genesegp-embl/AA1991.DAT:*	/SIDS1/gcgdata/hold-geneseg/genesegp-embl/AA1992.DAT:*	/SIDS1/gcgdata/hold-geneseq/genesegp-emb1/AA1993.DAT:*	/SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1994.DAT:*	/SIDS1/gcgdata/hold-geneseq/genesegp-emb1/AA1995.DAT:*	/SIDS1/gcgdata/hold-geneseq/genesegp-embl/AA1996.DAT:*	/SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1997.DAT:*	/SIDS1/9cgdata/hold-geneseq/genesegp-embl/AA1998.DAT:	/SIDS1/gcgdata/hold-geneseq/genesedp-embl/AA1999.DAT:*	/SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA2000.DAT:*	/SIDS1/gcgdata/hold-genesea/genesea/ashore
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	Rodent interleukin Mouse interleukin Murine IL-1 homolo Murine IL-1 recept Murine IL-ILI. Mu Invention related Murine interleukin Mouse interleukin Murine IL-1 recept A human interleukin
SUMMARIES	AAWB6284 AAY28407 AAY28260 AAY96937 AAB5261 AAB5261 AAB888 AAE06663 AAY97068 AAY97068
98	200 200 200 200 200 200 200 200 200 200
% Query Match Length DB	156 156 156 156 155 155 155 158 168 80
& Query Match	1000.0 1000.0 1000.0 99.4 99.4 99.4 99.4 99.4
Score	156 156 156 156 155 155 155 154 94
Result No.	1100 1100 1100

Protein encoded by Interleukin-II.1 re Human interleukin Human II.1 recepto Human II.1 homolog	IL-1 TANGO inter PRO43	eukin-IL1 n encoded eukin-IL1 human dia	Human IL-I recepto Interleukin ILI pe Generic human IL-I Hhuman IL-I homolo Generic human IL-I Epitope fragmen o	pitope fragment pitope fragment pitope fragment pitope fragment pitope fragment pitope fragment
AAB66663 AAB35263 AAY28408 AAY96936 AAY92256	AAY92257 AAY45062 AAE06655 AAB87601 AAB35260	AAB35262 AAB66664 AAB35264 ABG22711 AAY43526	AAY97067 AAB35266 AAY92253 AAY92254 AAY92255 AAW85942	AAW85941 AAW8594 AAW8594 AAW85943 AAW85944 AAW85945 AAW85945
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ALIGNMENTS

AAW86284 XX AAW86284 standard; Protein; 156 AA. AAW86284 standard; Protein; 156 AA. AC AAW86284; XX XX XX XX XX AAW86284; XX BT 19-FEB-1999 (first entry) BE Rodent interleukin (IL)-1 delta polypeptide. XX Interleukin; IL-1 delta; polyclonal antibody; IL-1 epsilon; cytokine; XX Interleukin; IL-1 delta; polyclonal antibody; IL-1 epsilon; cytokine; XX Interleukin; IL-1 delta; polyclonal antibody; IL-1 epsilon; cytokine; XX Interleukin; IL-1 delta; polyclonal antibody; IL-1 epsilon; cytokine; XX
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segment of IL-1 delta or IL-1 epsilon. Agonists or antagonists of these IL polypeptides are used to regulate a cell involved in an inflammatory response. The IL-1 delta or IL-1 epsilon polypeptides and peptides are used to produce Abs and antigen-Ab complexes. The polypeptides, Abs and the corresponding nucleic acids regulate development and/or the immune system, and can be used to diagnose and treat conditions associated with abnormal expression of IL. Agonists or antagonists of IL-1 delta or IL-1 epsilon polypeptides are used with agonists or antagonists of IL-1 delta or IL-1 LI-IRA, IL-1 betei, IL-1 gamma, IL-2 and/or IL-12. The IL-1 delta or IL-1 epsilon polypeptides may be used as a soluble polypeptide or as a fusion protein with another cytokine or chemokine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents mouse interleukin 1 delta (IL-1 delta). IL-1 delta proteins are useful for the determination of the molecular weight of a sample protein. The protein and its fragments are useful as controls for peptide fragmentation. This is useful for determining the
                                                                                                                                                                                                                                                                                                                                                                                                           61 GVQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 120
                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                            Interleukin 1 delta; IL-1 delta; glaucoma; ectodermal dysplasia;
insulin-dependent diabetes mellitus; wrinkly skin syndrome;
T-cell leukemia; lymphoma; tibial muscular dystrophy.
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                                                                                                                                                                                                                                                      100.0%; Score 156; DB 19;
100.0%; Pred. No. 5.6e-160;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY28407 standard; Protein; 156
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse interleukin 1 delta.
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Best Local Similarity 100.
Matches 156; Conservative
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                                                                                                                                                                                                           156 AA;
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09-JAN-1998;
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isoelectric point of a sample protein. Antibodies generated against CC iL-1 delta and its fragmented peptides can be used to enhance the accuracy of these molecular weight markers to determine the apparent concentration and isoelectric point of a sample protein. IL-1 delta can be used to screen for potential inhibitors of activity associated at therapeutic agents for the treatment of diseases amediated by IL-1 delta. IL-1 delta may be used as a reagent in studying the interleukin is called. IL-1 signalling pathway, or as a reagent to block IL-1 signalling. The IL-1 delta coding sequences can be used to identify human chromosome 2, and to identify genes associated with certain diseases, especially with region 2q11-12, including jaucoma, ectodermal dysplasia, insuling ceptable in muscular dystrophy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Proteins useful for treatment of inflammatory conditions such as rheumatoid arthritis and psoriasis are agonists or antagonists forms of new interleukin-1 homologue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Generic; interleukin-1; IL-1; homologue; zilla3; anti-inflammatory; antagonist; pro-inflammatory; agonist; immunomodulator; antiarthritic; antirheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic; immunosuppressive; antiulcer; antidiabetic; nephrotropic; Vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                  GVQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 120
                                                                                                                                                                                                                                                                                                                                                                                                                   1 MMVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVIL 60
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                                                                                                                                                                                                                                                                                 Length 156;
                                                                                                                                                                                                                                                                                 100.0%; Score 156; DB 20; 100.0%; Pred. No. 5.6e-160;
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                                                                                                                                                                                                                                                                                                                  Conservative
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N-PSDB; AAA09198.
                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                        156 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         vulnerary; 2q14
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                                                                                                                                                                                                                                                                                    Query Match
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used to analyze human northern blots.

It is believed that zilia3 acts through IL-1 receptors. In general, zilia3 proteins having a Lys residue at position 148 will have anti-inflammatory activity (e.g. AAY92256), whilst those having Asp (see AAY92254) or Glu at this position will have pro-inflammatory activity (e.g. AAY92256), whilst those having Asp (see AAY92254) or Glu at this position will have pro-inflammatory activity (e.g. AAY92256), whilst those having Asp (see AAY92254) or Glu at this position will have pro-inflammatory activity carrier or prevent chronic inflammatory diseases such as rheumatoid arthritis, osteoarthritis and Lyme arthritis, psoriasis, to reduce tissue damage after ischemia, to treat septic shock, graft-resuse host disease and leventae inflammatory bowel disease including Crohn's disease and ulcerative colitis, insulin-dependent disease mallitus, acute pancreatitis, glomerulonephritis and cerebral ischemia. Agonist forms of zilia3 may promote wound healing by IL-1 effects on growth factor secretion and cell proliferation. They may also treat infections, especially gastrointestinal infections.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TANGO-93; cytokine; mouse; secreted protein; IL-1 expression; cancer; Interleukin-1 receptor antagonist; IL-1ra; inflammation; antiasthmatic; immunosuppressive; antirheumatic; antiarthritic; antipsoriatic; asthma; antilnflammatory; antibacterial; antiulcer; cytostatic; immunomodulator; osteopathic; dermatological; antiulcer; cytostatic; immunomodulator; osteopathic; dermatological; antidiabetic; psoriasis; ulcerative colitis; graft vs-nost disease; rheumatoid arthritis; inflammatory bowel disease; septic shock; cachexia; crohn's disease; chronic myelogenous leukaemia; liver disease; diabetes; osteoarthritis; Hodgkin's disease; Lyme disease; autoimmune disease; myasthenia gravis; pharmacogenomic; diagnosis; systemic lupus erythematosus; forensic; transgenic animal.
                      is shows an interleukin-1 (IL-1) homologue, designated zilla3. A 350 probe generated from the DNA sequence by PCR using AAA09199-200 was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61 GVQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 120
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Best Local Similarity 100.
Matches 156; Conservative
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belongs to the cytokine superfamily. It plays a role similar to secreted profesh, the cytokine superfamily. It plays a role similar to secreted for the cytokine superfamily. It plays a role similar to secreted interleukin-1 receptor antagonist (IL-1ra) and its expression is developmentally regulated in liver, heart and bone marrow. TangO-93 modulates immune mediated inflammation and IL-1 gene or protein expression. TangO-93 is useful as a modulating agent for regulating cellular processes like asthma, graft vs-host disease, rhemmatoid arthritis, psortasis, inflammatory bowel disease, septic shock, ulcerative colitis, Crohn's disease, chronic myelogenous leukaemia, cancer, liver disease, Hodokin's disease, osteoarthritis, Lyme disease, cachexia, and autoimmune diseases or, myasthenia gravis, autoimmune diabetes and systemic lupus erythematosus. Partial TangO-93 sequences are useful in forensic blology, for diagnostic and prognostic assays, prophylactic and therapeutic treatment and pharmacogenomics. The DNA sequences are useful as hybridisation probes and primers, for isolation of TangO-93 sequence and for the creation of transgenic animals.
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                                                                                                                                                                                                    The present sequence is the murine TANGO-93, a secreted protein, that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61 GVQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated nucleic acid sequences encoding TANGO-93 polypeptide useful for treating a variety of cellular processes e.g. asthma, rheumatoid arthritis, psoriasis and autoimmune diseases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mIL-1Ra3; interleukin-1 receptor antagonist-3; IL-11p; osteopathic; interleukin-1-like polypeptide; anti-inflammatory; anti-asthmatic; anti-arthritic; antimicrobial; respiratory; anti-ischemic; vaccine; dermatalogical; immunomodulatory; gastrointestinal; gene therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 156;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 156; DB 21;
100.0%; Pred. No. 5.6e-160;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               121 FLCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Murine IL-1 receptor antagonist 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY96937 standard; Protein; 155
                                                                                                                                                                   Claim 9; Fig 1; 113pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99WO-US30720.
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Best Local Similarity 100.1
Matches 156; Conservative
                                     WPI; 2000-205669/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            156 AA;
                                                       N-PSDB; AAZ50811
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22-JAN-1999;
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(HYSE-) HYSEQ INC.
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ID AAB6
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                                                                                                                                                                                         polypeptide (11-11p) that retains one or more activities of the peptide from which it is derived, such as the IL-18R binding activity of a human interleukin-1 receptor antagonist-1 (hIL-1Ral) polypeptide, is new. The nucleic acids may be used in molecular engineering applications, e.g. hybridization assays and chromosome and gene mapping studies, for recombinantly producing the IL-11p polypeptide or for producing gene knock out animals to study the role of the protein in metabolism and disease processes (conversely, gene therapy protecols may be used to supplement a pathents production of the polypeptide or to rectify mutations that lead to the production of in active peptides). The peptides produced may be used to screen for and produce modulators (e.g.
                                                                                                                                                                                                                                                                                                                                   antibodies) of IL-IIP protein expression and activity which may be use
to treat disorders associated with inappropriate IL-IIP expression and
activity, such as inflammatory disorders, asthma, arthritis,
osteoarthritis, sepsis, acute lung injury, adult respiratory distress
syndrome, idiopathic pulmonary fibrosis, ischemic reperfusion disease,
psoriasis, graft versus host disease and/or inflammatory bowel disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse; IL-IL1; interleukin-1 locus; IL-1beta; IL-1receptor; psoriasis; chromosome 2013; inflammatory disease; heart disease; Graves' disease; rheumatoid arthritis; inflammatory bowel disorder; diabetes; cancer; osteoporosis; systemic lupus erythematosus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 vqqqsqclscgtekgpilklepvnlmelylqakesksftfyrrdmgltssfesaaypgwf 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       62 VQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRDMGLTSSFESAAYPGWF 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILG 61
                                                                                                                                                                                 isolated nucleic acid molecule encoding an interleukin-1-like
                                                                                                                   Nucleic acids encoding interleukin-1-like polypeptides, useful i
preventing and treating e.g. inflammation, asthma and psoriasis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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6.7e-159;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99.4%; Score 155; DB
100.0%; Pred. No. 6.7
:ive 0; Mismatches
                                                                                                                                                         Claim 22; Fig 9A-B; 143pp; English.
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 99US-0129122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                         (GETH ) GENENTECH INC
                                                                             2000-452395/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 155; Conserv
                                                    Pan J;
                                                                                                                                                                                                                                                                                                                                                                                                                                155 AA;
                                                                             WPI; 2000-452395,
N-PSDB; AAA51599
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13-APR-1999;
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                                                   Goddard A,
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AAB35261
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The present invention provides the protein and coding sequences of the human and murine interleukin-ILI (IL-ILI) proteins. The IL-ILI gene is located between the IL-Ibeta and IL-Ireceptor genes at human chromosome 2q13. The sequences are useful in the diagnosis, prevention and treatment of heart disease, cancer and inflammatory diseases such as rheumatoid arthritis, systemic lupus erythematosus, inflammatory bowel disorder, diabetes, psoriasis, osteoporosis, lichen sclerosis, ulcerative colitis, severe periodontal disease and pregnancy complications. The present sequence is the murine IL-ILI protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       62 VQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61
                                                                                                                                                                                                                                                                                       Nucleic acids encoding human and murine interleukin-1L1 polypeptides useful for controlling inflammatory processes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 MYLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 99.4%; Score 155; DB 22; I Best Local Similarity 100.0%; Pred. No. 6.7e-159; Matches 155; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       121 lctspeadqpvrltqipedpawdapitdfyfqqcd 155
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                                                                                                                   (INTE-) INTERLEUKIN GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                          Claim 11; Fig 3; 150pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Invention related sequence #4.
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990S-0457626.
2000US-0523552.
2000US-0576008.
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                                                           99US-0144298
17-JUL-2000; 2000WO-US19508
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                                                                                                                                                                            Barton J;
                                                                                                                                                                                                                                 WPI; 2001-091974/10.
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08-DEC-1999;
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22-MAY-2000;
                                                           16-JUL-1999;
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09-AUG-2001
       Smith RF,
                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Interleukin-1 homologue 3; IL-1H3; mouse; murine; drug screening; agonist; antagonist; human disease; chronic inflammation; acute inflammation; septicaemia; autoimmune disease; psoriasis; inflammatory bowel disease; arthritis; transplant rejection; infection; graft versus host disease; acute respiratory distress syndrome; allergy; asthma; restenosis; stroke; ischaemia; brain injury; AIDS; bone disease; concer; lymphoporoliferative disorder; atherosclerosis; congestive heart failure; Alzheimer's disease; immunosuppressive;
                                                  Isolated nucleic acids encoding interleukin-1 (IL-1) receptor antagonist proteins (referred as IL-1Hyl), useful in the treatment of cancer, e.g. breast adenocarcinoma and brain tumors, and an inflammatory disease mediated by IL-18 -
                                                                                                                                                                                                                                                                                                                              VQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 121
                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                     2 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILG
                                                                                                                       The present invention relates to interleukin (IL)-1 receptor antegonist proteins. IL-1Hyl is useful for treating cancer, an inflammatory disease mediated by IL-18, inflammation resulting from infection or allergic reactions, and inflammation associated with chronic bronchitis, arthritis, diabetes or
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                                                                                                                                                                                                                                     Length 155;
                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                 99.4%; Score 155; DB 22; I
100.0%; Pred. No. 6.7e-159;
Live 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Murine interleukin-1 homologue 3 (IL-1H3).
                                                                                                     Disclosure; Page 177; 179pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB48828 standard; Protein; 155 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antimicrobial; neuroprotective
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2000US-0577715.
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                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.
Matches 155; Conservative
                              WPI; 2001-071582/08.
                                                                                                                                                                                                   Sequence 155 AA;
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23-MAY-2000;
                                                                                                                                                                               endothermia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-NOV-2000.
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         Ford J,
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AAB48828
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The invention also relates to expression vectors and host calls comprising murine IL-IH3 nucleic acids, the recombinant production of murine IL-IH3 and one of screening for medilators of IL-IH3 agonists and antagonists thus identified. IL-IH3 agonists and antagonists thus identified a IL-IH3 agonists and acute inflammation, septicaemia, autoimmune diseases (e.g., inflammatory bowel disease, infection, stroke, ischaemia, rejection, graft versus host disease, infection, stroke, ischaemia, brain injury, AIDS, bone diseases (e.g., osteoporosis), cancer setting injury, AIDS, bone diseases (e.g., osteoporosis), cancer atheroscierosis, and Alzheimer's disease. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse; interleukin-1delta; IL-1delta; virucide; hepatotropic; fever; immunological disorder; tumour; inflammatory disorder; hypoglycaemia; autoimmune disease; pulmonary tuberculosis; fulminant hepatitis; leprosy; psoriasis; viral infection; allergy; cytokine; HIV; drug screening.
                                                                                                                                                                                                                                                                                                                    The invention relates to murine interleukin-1 homologue 3 (IL-1H3; AAB48828) and nucleic acids which encode it (cDNA given in AAC81700), including nucleic acid sequences with at least 95% identity to AAC81700.
                                                                                                                                                  used for screening modulators treating autoimmune diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              62 VQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 121
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Pred. No. 6.7e-159;
   Halsey W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse interleukin-1delta (IL-1delta) protein.
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                                                                                                                                         Murine interleukin-1 homolog polypeptide of the polypeptide which can be used for cancer, brain injury and bone disorders
McDonnell PC,
                                                                                                                                                                                                                                                         Claim 2; Page 28-29; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAE06663 standard; Protein; 154 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99.4%; SCC.
100.0%; Pre
0; }
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Best Local Similarity 100.
Matches 155; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              represents murine IL-1H3
Young PR,
                                                    2001-025138/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       155 AA;
                                                                                 N-PSDB; AAC81700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200157219-A2.
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                                                                                                                                                                      The invention relates to recombinant antigenic interleukin-1 like molecules and their corresponding nucleic acid sequences, designated as interleukin-idelita [IL-idelta] and interleukin-idelita]. [IL-idelta] and interleukin-idelta] [IL-idelta] and interleukin such as immunological disorders, tundurs, inflammatory disorders, fever, hypoglycaemia, psoriasis, tumborus, inflammatory disorders, fever, hypoglycaemia, psoriasis, tumborus, inflammatory disorders, fever, hypoglycaemia, psoriasis, tumborus, leaven inflammatory disorders, fever, hypoglycaemia, psoriasis, tumborus, leaven inflaminant hepatitis, and viral infections such as HIV. The invention also relates to methods of using the composition containing IL-idelta or IL-idepsilon for both diagnostic and therapeutic utilities. IL-idelta is used as an immunogen for the production of antisera or antibodies specific, e.g., capable of distinguishing between inspection or immunological disorders which lead to antibody production captus or immunological disorders which lead to antibody production competitive drug screening assays. The presence of elevated levels of competitive drug screening assays. The presence is mouse of competitive drug screening assays. The presence is mouse of interleukin-idelta (IL-idelta) protein related to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                              Novel isolated or recombinant antigenic interleukin-1 delta or epsilon polypeptide useful for treating conditions exhibiting abnormal expression of interleukin such as immunological disorders, tumor and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             63 QGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFL 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 VLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGV 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mIL-1Ra3; interleukin-1 receptor antagonist-3; IL-11p; osteopathic; interleukin-1-1like polypeptide; anti-infilammatory; anti-asthmatic; anti-arthritic; antimicrobial; respiratory; anti-ischamic; vaccine; dermatalogical; immunomodulatory; gastrointestinal; gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98.7%; Score 154; DB 22; Length 154; 100.0%; Pred. No. 8e-158; Live 0; Mismatches 0; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Murine IL-1 receptor antagonist 3 EST W08205 product.
                           Kastelein RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  123 CTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bazan JF,
                                                                                                                                                    Disclosure; Fig 1; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY97068 standard; Protein; 94 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.
Matches 154; Conservative
                          Timans JC,
(SCHE ) SCHERING CORP.
                                                    WPI; 2001-488886/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          154 AA;
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                           Debets JEMA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-JUL-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                           allergy
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polypeptide (IL-IIP) that retains one or more activities of the peptide from which it is derived, such as the IL-IBR binding activity of a human interleukin-1 receptor antagonist-1 (ILI-IRA1) polypeptide. Is new. The nucleic acids may be used in molecular engineering applications, e.g. hybridization assays and chromosome and gene mapping studies, for recombinantly producing the IL-IIP polypeptide or for producing gene knock out animals to study the role of the protein in metabolism and disease processes (conversely, gene therapy protocols may be used to supplement a patients production of the polypeptide or to rectify mutations that lead to the production of in active peptides). The peptides produced may be used to screen for and produce modulators (e.g. antibodies) of IL-IIP protein expression and activity which may be use to treat disorders associated with inappropriate IL-IIP expression and activity, such as inflammatory disorders, asthma, arthritis, sepsis, acute lung injury, adult respiratory distress syndrome, idiopathic pulmonary fibrosis, ischemic reperfusion disease. Sociassis, graft versus host disease and/or inflammatory bowel disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            An isolated nucleic acid molecule encoding an interleukin-1-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acids encoding interleukin-1-like polypeptides, useful f
preventing and treating e.g. inflammation, asthma and psoriasis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 94;
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2.5e-93;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           60.3%; Score 94; DB 100.0%; Pred. No. 2.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 gvgggsgclscgtekgpilklepvnimelylgak
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Fig 10; 143pp; English.
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                                                             98US-0113430.
99US-0116843.
99US-0129122.
99WO-US30720
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                                                                                                                                                                                                           GETH ) GENENTECH INC
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N-PSDB; AAA51600.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94 AA;
22-DEC-1999;
                                                                     23-DEC-1998
                                                                                                                                      13-APR-1999
                                                                                                     2-JAN-1999
                                                                                                                                                                                                                                                                             Goddard A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY43525;
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Protein encoded by B2HFLS20W cDNA library sequence #2.

interleukin; IL-1 receptor; cancer; inflammation

99US-0348942. 99US-0417455. 99US-0457626. 2000US-0523552.

13-OCT-1999; 08-DEC-1999; 10-MAR-2000;

07-JUL-1999;

22-MAY-2000;

WPI; 2001-071582/08.

Pace A;

Ford J,

(HYSE-) HYSEQ INC.

2000WO-US18710

07-JUL-2000;

11-JAN-2001

WO200102571-A2.

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The present sequence represents a human interleukin-1 (IL-1) receptor antagonist. It is encoded by cDNA sequences obtained from the b2HFLS20W cDNA library of foetal liver-spleen. The polypeptide is capable of binding IL-1 receptors (IL-1Rs). The polynuclectides and polypeptides can binding IL-1 receptors (IL-1Rs). The polynuclectides and polypeptides can be used for the prevention or treatment of disorders involving sepsis, caute pencreatitis, endotoxic shock, cytokine induced shock, rheumatoid arthritis, chronic inflammatory arthritis, pancreatic cell damage from disease winflammatory disease, inflammatory bowel cause or inflammatory disease, inflammatory disease, an antiproliferative agent such as for acute or chronic myelogenous leukemia or in the prevention of premature caute or chronic myelogenous leukemia or in the prevention of premature cautivities such as e.g. nutritional activity, immune stimulating or proliferation/differentiation activity, immune stimulating or suppressing activity, hematopoiesis regulating activity, tissue growth emostatic and thrombolytic activity, receptor/ligand activity, and activity, activity, receptor/ligand activity, and anti-inflammatory activity. The products can also be used for contents of the products can also be used for the contents.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated interleukin-1 receptor binding polypeptides, used to treat e.g. sepsis, shock, arthritis, pancreatitis, graft-versus-host disease, inflammatory disease, autoimmune disease or proliferative disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
activin activity; inhibin activity; chemotactic activity;
chemokinetic activity; hemostatic activity; thrombolytic activity;
anti-inflammatory activity.
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Alfenito M;
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                                                                                                                                                                                                                                                                                                                                                                                                                   Dickson M, Dr
d J, Pace A,
                                                                                                                                                                                                                                                                                                                                                                                                                 Crkvenjakov R, Dick
J. Kita D, Ford J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 8; Fig 3; 123pp; English.
                                                                                                                                                                                                                                               98US-0079909
98US-0082364.
98US-0099818.
98US-0127698.
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                                                                                                                                                                                                                                                                                                                                                                                 (HYSE-) HYSEQ INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    80
                                                                            Homo sapiens.
                                                                                                            W09951744-A2
                                                                                                                                                                                        05-APR-1999;
                                                                                                                                                                                                                                                                                                                         13-JAN-1999;
17-FEB-1999;
                                                                                                                                                    14-OCT-1999
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                                                                                                                                                                                                                                                                   20-MAY-1998
                                                                                                                                                                                                                                                                                                         31-JUL-1998
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Matches 4
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Isolated nucleic acids encoding interleukin-1 (IL-1) receptor antagonist proteins (referred as IL-1Hy1), useful in the treatment of cancer, e.g. breast adenocarcinoma and brain tumors, and an inflammatory disease mediated by IL-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse; IL-IL1; interleukin-1 locus; IL-lbeta; IL-lreceptor; psoriasis; chromosome 2d13; inflammatory disease; heart disease; Graves' disease; rheumatoid arthritis; inflammatory bowel disorder; diabetes; cancer; osteoporosis; systemic lupus erythematosus; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to interleukin (IL)-1 receptor antigonist proteins. IL-1Hy1 is useful for treating cancer, an inflammatory disease mediated by IL-18, inflammation resulting from infection or allergic reactions, and inflammation associated with chronic bronchitis, arthritis, diabetes or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28.2%; Score 44; DB 22; I 100.0%; Pred. No. 1.7e-39; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Interleukin-IL1 recombinant protein #2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB35263 standard; Protein; 154
                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Fig 3; 179pp; English.
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Best Local Similarity
Matches 44; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          80 AA;
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AAB66663 standard; protein;

12

RESULT 1

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(first entry)

05-APR-2001

AAB66663;

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AAY96936;
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Matches
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                                                                                                                               human and murine interjeukin-ILI (ILTII) proteins. The IL-ILI gene is located between the IL-lbeta and IL-Irceptor genes at human chromosome 2q13. The sequences are useful in the diagnosis, prevention and treatment of heart disease, cancer and inflammatory diseases such as rheumatoid arthritis, systemic lupus erythematosus, inflammatory bowel disorder, severe periodontal disease and pregnancy complications. The present sequence is a recombinant IL-ILI protein.
                                                                                                                        The present invention provides the protein and coding sequences of the
                                                                                                                                                                                                                                                        Gaps
                                                                             human and murine interleukin-1L1 polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                      Interleukin 1 delta; IL-1 delta; glaucoma; ectodermal dysplasia; insulin-dependent diabetes mellitus; wrinkly skin syndrome; T-cell leukemia; lymphoma; tibial muscular dystrophy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Murine and Human interleukin 1 delta DNA, polypeptides and its fragments, useful as molecular weight markers
                                                                                                                                                                                                                                                        ;
0
                                                                                                                                                                                                                                       Length 154;
                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                  81 LEPVNIMELYLGAKESKSFTFYRDMGLTSSFESAAYPGWFLCT 124
                                                                                                                                                                                                                                      28.2%; Score 44; DB 22; I
100.0%; Pred. No. 3.3e-39;
ive 0; Mismatches 0;
                                                                                      inflammatory processes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 68-69; 72pp; English
                                                                                                                                                                                                                                                                                                                                     AAY28408 standard; Protein; 155 AA
                        (INTE-) INTERLEUKIN GENETICS INC.
                                                                                                        Examples; Fig 6; 150pp; English.
         99US-0144298
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                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                         Human interleukin 1 delta.
                                                                                                                                                                                                                                                        44; Conservative
                                                                             Nucleic acids encoding useful for controlling
                                           Nicklin M, Barton J;
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                                                            WPI; 2001-091974/10.
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                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                              154 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB; AAX89432.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
         16-JUL-1999;
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                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                        Matches
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The present sequence represents human interleukin 1 delta (IL-1 delta).

IL-1 delta proteins are useful for the determination of the molecular weight of a sample protein. The protein and its fragments are useful as controls for peptide fragmentation. This is useful for determining the isoelectric point of a sample protein. Antibodies generated against IL-1 delta and its fragmented peptides can be used to enhance the accuracy of these molecular weight markers to determine the apparent molecular eight and isoelectric point of a sample protein. IL-1 delta can be used to screen for potential inhibitors of activity associated with IL-1 delta counter-structure molecules. IL-1 delta can also be used as therapeutic agents for the treatment of diseases mediated by IL-1 delta in IL-1 delta may be used as a reagent in studying the interleukin I can identify genes associated with certain diseases, especially with region 2dil-12, including glaucoma, ectodermal dysplasia, insulin-debendent diabetes mellitus, wrinkly skin syndrome, T-cell leukemia/ lymphoma and tibial muscular dystrophy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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100.0%; Pred. No. 3.3e-39;
:Ive 0; Mismatches 0;
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99US-0116843.
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Best Local Similarity
Matches 44; Conserv
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13-APR-1999;
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interleukin-1 receptor antagonist-1 (hIL-IRal) polypeptide, is new. The nucleic acids may be used in molecular engineering applications, e.g. hybridization assays and chromosome and gene mapping studies, for recombinantly producing the IL-Ilp polypeptide or for producing gene knock out animals to study the IL-Ilp polypeptide or for producing gene disase processes (conversely, gene therapy protocols may be used to supplement a patients production of the polypeptide or to rectify mutations that lead to the production of the polypeptide or to rectify peptides produced may be used to supplement a patients production of the polypeptide or to rectify netations that lead to the production of in active peptides). The peptides produced may be used to screen for and produce modulators (e.g. antibodies) of IL-Ilp protein expression and activity which may be use to treat disorders associated with inappropriate IL-Ilp expression and activity, which may be use contractly distrementation, sepsis, acute lung injury, adult respiratory distress syndrome, idiopathic pulmonary fibrosis, ischemic reperfusion disease.
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155 AA; Sequence

; 0 0; Gaps Query Match 28.2%; Score 44; DB 21; Length 155; Best Local Similarity 100.0%; Pred. No. 3.3e-39; Matches 44; Conservative 0; Mismatches 0; Indels 0; Indels

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Search completed: May 22, 2002, 16:18:25 Job time: 121 sec

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                                                                                                             May 22, 2002, 16:16:39; Search time 13.05 Seconds
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Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                     OM protein - protein search, using sw model
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Gapop 60.0 , Gapext 60.0
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                                                                                                                                                                                                                                 Sequence:
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/cgn2_6/ptodata/2/laa/5B_COMB.pep:*
/cgn2_6/ptodata/2/laa/6A_COMB.pep:*
/cgn2_6/ptodata/2/laa/6B_COMB.pep:*
/cgn2_6/ptodata/2/laa/RCOMB.pep:*
/cgn2_6/ptodata/2/laa/PCTUS_COMB.pep:* Issued_Patents_AA:* Database :

Post-processing: Listing first 1000 summaries

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES Query Match Length DB Score

Sequence 3, Sequence 5, Sequence 5, Sequence 5, Sequence 5, Description US-09-417-455-3 US-09-348-942-3 US-09-417-455-5 US-09-348-942-5 US-09-316-081-5 28.2 28.2 28.2 28.2 28.2

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ALIGNMENTS

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Sequence 3, Application US/09417455
Patent No. 6294655
Patent Pack, Ann
APPLICANT: Pace, Ann
APPLICANTON: A NOVEL INTERLEUKIN-1 RECEPTOR ANTAGONIST AND USES THEREOF
FILE REFERENCE: 28110/36328
CURRENT APPLICATION NUMBER: US 09/348,942
PRIOR APPLICATION NUMBER: PGT/US99/04291
PRIOR FILING DATE: 1999-07-07
PRIOR FILING DATE: 1999-04-05
PRIOR APPLICATION UNBER: US 09/287,210
PRIOR APPLICATION UNBER: US 09/287,210
US-09-417-455-3
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Sequence 3, Application US/09348942

Sequence 3, Application US/09348942

Sequence 3, Application US/09348942

GENERAL INCPRANTION:

FILE OF INVENTION: A NOVEL INTERLEUKIN-1 RECEPTOR ANTAGONIST AND USES THEREOF

CURRENT APPLICATION NUMBER: US/09/348,942

CURRENT APPLICATION NUMBER: US/09/348,942

CURRENT FILING DATE: 1999-07-07

EARLIER APPLICATION NUMBER: US/09/287,210

EARLIER APPLICATION NUMBER: US/09/281,370

EARLIER PILING DATE: 1999-04-05

EARLIER PILING DATE: 1999-01-13

EARLIER PILING DATE: 1999-01-13

EARLIER PILING DATE: 1999-07-31

EARLIER PILING DATE: 1999-07-31

EARLIER PILING DATE: 1998-06-19

EARLIER PILING DATE: 1998-06-19
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28.2%; Score 44; DB 4; L
Best Local Similarity 100.0%; Pred. No. 2.2e-38;
Matches 44; Conservative 0; Mismatches 0;
PRIOR APPLICATION NUMBER: US 09/251,370
PRIOR FILING DATE: 1999-02-17
PRIOR PILING DATE: 1999-01-13
PRIOR FILING DATE: 1999-01-13
PRIOR PLING DATE: 1998-01-31
PRIOR APPLICATION NUMBER: US 09/127,698
PRIOR PILING DATE: 1998-06-19
PRIOR PILING DATE: 1998-06-19
PRIOR PILING DATE: 1998-06-19
PRIOR PILING DATE: 1998-05-20
PRIOR PILING DATE: 1998-05-20
PRIOR PILING DATE: 1998-05-15
PRIOR PILING DATE: 1998-05-15
PRIOR PILING DATE: 1998-06-15
PRIOR PILING DATE: 1998-04-03
NUMBER OF SEQ ID NOS: 30
SECTRARRE: FRASESE FOR WINDOWS VEISION 3.0
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 3
LENGTH: 80
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; ORGANISM: Homo sapiens
US-09-417-455-3
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Matches 4½; Conserva
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Gaps

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Length 155; Indels

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APPLICANT: Ballinger, Dennis G.
APPLICANT: Ballinger, Dennis G.
APPLICANT: Bace, Ann M.
TITLE OF INVENTION: Interleukin-1 Hy2 Materials and Methods
FILE REPERENCE: 28110/35659
CURRENT APPLICATION NUMBER: US/09/316,081
CURRENT FILING DATE: 1999-05-20
NUMBER OF SEO ID NOS: 11
SOFTWARE: Patentin Ver. 2.0
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Pred. No. 3.9e-38;
0; Mismatches 0;
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28.2%; Score 44; DB 4; L
Best Local Similarity 100.0%; Pred. No. 3.9e-38;
Matches 44; Conservative 0; Mismatches 0;
     EARLIER APPLICATION NUMBER: US 09/099,818
EARLIER FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: US 09/082,364
EARLIER APPLICATION NUMBER: US 09/079,909
EARLIER APPLICATION NUMBER: US 09/079,909
EARLIER FILING DATE: 1998-05-15
EARLIER FILING DATE: 1998-07-03
SOFTWARE: PASTSEQ for Windows Version 3.0
SSOFTWARE: FASTSEQ for Windows Version 3.0
EENGTH: 155
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Job time: 125 sec
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ilarity 100.0%; Pr
Conservative 0;
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US-09-316-081-5
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US-09-348-942-5
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Best Local Similarity
Matches 44; Conserv
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TYPE: PRT
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APPLICANT: Ford, John
APPLICANT: Ford, John
TITLE OF INVERIL INCREDENT A NOVEL INTERLEUKIN-1 RECEPTOR ANTAGONIST AND USES THEREOF
FILE REFERENCE: 28110/36328
CURRENT PAPLICATION NUMBER: US/09/417,455
CURRENT FILING DATE: 1999-10-13
PRIOR PAPLICATION NUMBER: US 09/348,942
PRIOR PAPLICATION NUMBER: PCT/US99/04291
PRIOR PAPLICATION NUMBER: PCT/US99/04291
PRIOR PLING DATE: 1999-04-05
PRIOR APPLICATION NUMBER: US 09/251,370
PRIOR PAPLICATION NUMBER: US 09/251,370
PRIOR PAPLICATION NUMBER: US 09/251,370
PRIOR PLING DATE: 1999-01-13
PRIOR PLING DATE: 1999-01-13
PRIOR PLING DATE: 1999-01-31
PRIOR PLING DATE: 1999-01-31
PRIOR PLING DATE: 1998-05-15
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TITLE OF INVENTION: A NOVEL INTERLEUKIN-1 RECEPTOR ANTAGONIST AND USES THEREOF

FILE REFERENCE: 28110/35801

CURRENT APPLICATION NUMBER: US/09/348,942

CURRENT FILING DATE: 1999-07-07

EARLIER APPLICATION NUMBER: CT/US99/04291

EARLIER PILING DATE: 1999-04-05

EARLIER PLING DATE: 1999-04-05

EARLIER PILING DATE: 1999-04-05

EARLIER PILING DATE: 1999-04-05

EARLIER PILING DATE: 1999-02-17
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81 LEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCT 124
                              Query Match 28.2%; Score 44; DB 4; Le Best Local Similarity 100.0%; Pred. No. 3.9e-38; Matches 44; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 1999-02-17
APPLICATION NUMBER: US 09/229,591
FILING DATE: 1999-01-13
APPLICATION NUMBER: US 09/127,698
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 5, Application US/09348942 Patent No. 6337072
                                                                                                                               RESULT 3
US-09-417-455-5
Sequence 5. Application US/09417455
Fatent No. 6294655
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: Homo sapiens
US-09-417-455-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Fast
SEQ ID NO 5
LENGTH: 155
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RC STRAINCSTBL/65; TISSUE-TONGUE, AND STOMACH;
RX MANI J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,
RX MAWI J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,
RX Arakawa T., Hara A., Eukunishi Y., Konno H., Adachi J., Fukuda S.,
RX Aito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
Rajto T., Okasaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
Radito T., Okasaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
Radota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Schriml L.M., Eaubli F., Suzuki R., Caninci P., de Bonaldo M.F.,
RA Schriml L.M., Bolunga N., Carninci P., de Bonaldo M.F.,
RA Sakai K., Okido T., Fletcher C., Fulta M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Wishaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
RA Wanshar-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
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                                                                                        Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Functional annotation of a full-length mouse cDNA collection.";
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156 AA: 17136 MW; A4D1EE2F93CF77A7 CRC64;
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090YY1;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 156; DB 11;
100.0%; Pred. No. 1.5e-158;
tive 0; Mismatches 0;
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Interpro, IPR000975; Interleukin_1.
ProDom; PD002536; Interleukin_1; 1.
SMART; SM00125; ILI; 1.
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EMBL; AF230378; AAF91275.1; -.
EMBL; AK009741; BAB26471.1; -.
EMBL; AK008977; BAB26002.1; -.
HSSP; P18510; 11LR.
MGD; MGI:1859325; 111f5.
                                                                                     Eukaryota; Metazoa; Chordata;
Mammalla; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                            Mus musculus (Mouse).
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Best Local Similarity
Matches 156; Conserv
                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                              NCBI_TaxID=10090;
                              OR ILIHY1.
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SEQUENCE
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                                                                                                                                                                       May 22, 2002, 16:18:30 ; Search time 27.05 Seconds (Without alignments) 997.679 Million cell updates/sec
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Q9qyy1 mus musculu
Q9ubh0 homo sapien
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156
1 MMVLSGALCFRMKDSALKVL.....IPEDPAWDAPITDFYFQQCD 156
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01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEUKIN-1 DELTA (INTERLEUKIN 1 RECEPTOR ANTAGONIST HOMOLOG
            GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Gapop 60.0 , Gapext 60.0
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Q9UBH0
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sp_unclassified:*
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sp_human:*
sp_invertebrate:*
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sp_organelle:*
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Maximum DB seq length: 200000000
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: sp_bacteria:*
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RESULT **09JIG2**

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Search completed: May 22, 2002, 16:21:51
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Matches 44;
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                                                                                                                                                                                                                                                                                                                                                                                                Identification and initial characterization of four novel members
                                                                                                                                                                                                                                                                                                           MEDLINE-20209405; PubMed-10744718;
Kumar S., McDonnell P.C., Lehr R., Tierney L., Tzimas M.N.,
Griswold D.E., Capper E.A., Tal-Singer R., Wells G.I., Doyle M.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
FILL DELTA (INTERLEUGHIN-1 LIKE PROTEIN 1) (INTERLEUKIN-1 RECEPTOR ANTAGONIST HOMOLOG 1) (INTERLEUKIN-1 DELTA).
                                                                Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Euteleostomi;
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                                                                                                                                                                        Barton J.L., Nicklin M.J.H.;
"ILL-ILI: A Novel Member of the Interleukin-1 Gene Family is
in Trophoblasts and Macrophages.";
Submitted (OCT-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 155;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prodom; PDU02350, ILI; 1.
SMART; SM00125; ILI; 1.
PROSTIE; PS00253; INTERLEUKIN_1; 1.
155 AA; 17004 MW; A4B1770F2E12533A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Butheria; Primates; Catarrhini; Hominidae;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99.4%; Score 155; DB 11; I 100.0%; Pred. No. 1.7e-157; ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Smith D.E., Renshaw B.R., Ketchem R.R., Kubin M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    *Four New Members Expand the IL-1 Superfamily.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 155 AA.
  PROTEIN (INTERLEUKIN-1 HOMOLOG 3).
                                                                                                                                                                                                                                                                                                                                                                                                                   the interleukin-1 family.";
J. Biol. Chem. 275:10308-10314(2000).
EMBL, AP250495; CAB59831.1;
EMBL, AP20495; AAF69251.1;
HSSP; P18510; JILR.
MGD; MGT:1859325; JILFS.
InterProc; IPPR000975; Interleukin_1.
ProDom, PD002536; Interleukin_1; 1.
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                                           Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Best Local Similarity 100.
Matches 155; Conservative
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SEQUENCE FROM N.A.
                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                         NCBI_TaxID=10090;
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  IL-1L1 PROTEIN IL1F5 OR IL1HY1
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Barton J.L., Herbst R., Bosisio D., Nicklin M.J.H.;
"A tissue specific interleukin-1 receptor antagonist homolog from the IL-1 cluster lacks IL-1. IL-1ra, IL-18 and IL-18ra activities."; submitted (JAN-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                from the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Novel IL-1 ligands IL-1d and IL-1e use IL-1R related protein 2."; Submitted (FEB-2000) to the EMBL/GenBank/DDBJ databases. EMBL; AF201830; AAF25210.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        [5]
SEQUENCE FROM N.A.
MEDLINE-20322477; PubMed-10866108;
Mulero J.J., Nelken S.T., Ford J.E.;
"Organization of the Human Interleukin-1 Receptor Antagonist Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                              Barton J.L., di Giovine F.S., Symons J.A., Nicklin M.J.H.;
"A tissue specific interleukin-1 receptor antagonist homolog i
ILI cluster lacks IL-1, IL-1ra, IL-18 and IL-18ra activities.'
Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.
MEDLINE-99443727; PubMed-10512743;
Mulero J.J., Pace A.M., Nelken S.T., Loeb D.B., Correa T.R.,
Drmance R., Ford J.E.;
"ILLHY1: A Novel Inter:
Biochem. Biophys. Res. Commun. 263:702-706(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 155;
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Debets R., Timans J.C., Zurawski S., Sana T.R., Bazan F.,
Kastelein R.A.;
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8e-39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROSITE; PS00253; INTERLEUKIN_1; UNKNOWN_1
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Pred. No.
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ilarity 100.0%; Pr
Conservative 0;
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EMBL, AJ242738; CAB59833.1;
EMBL, AJ271338; CAB67704.1;
EMBL, AF21693; ARF76981.1;
EMBL, AF230377; AAF91274.1;
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nes 44; Conserv
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May 22, 2002, 14:05:33 ; Search time 29.87 Seconds (without alignments) 580.097 Million cell updates/sec
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1 MMVLSGALCFRMKDSALKVL.....IPEDPAWDAPITDFYFQQCD 156
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                 OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Perfect score:
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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Minimum DB seq length: 0 Maximum DB seq length: 200000000

A_Geneseq_032802:* Database :

1. (SIDSI)/gcgdata/hold geneseq/geneseqp-embl/AA1980.DAT: *
2. (SIDSI/gcgdata/hold geneseq/geneseqp-embl/AA1981.DAT: *
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20. (SIDSI/gcg /SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1999.DAT:*/SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA2000.DAT:*/SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA2001.DAT:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		Description		Rodent interleukin	Money intonionitie	Munice Incelledail	MULTING IL-I DOMOIO	Murine TANGO-93 pr	Muripo II -1 2000	Ministra II I Coppe	Murine IL-ILI. Mu	Invention related	TOTAL	war the three teath	Mouse interleukin-	Transporter, demand	unwan Threftenkin	' Human IL-1 recepto
SUMMARIES		ΩI	700000000	PG700MW	AAY28407	AAV92260		AAY45061	AAY96937	AAB35261	TOTOCOM	AAB66672	AABABAB		AAEU6663	AAY28408		AAY 96936
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Human IL-1 homolog	Human TANGO-93 pro	interlenk		IL-ILL H	Ξ		interlen	Interleukin-IL1 re		-	Hhuman II1		human	Novel human diagno	Murine II			Protein encoded by	- 0	Interleukin-1 rece	Mouse interleukin-			Mouse interleukin-		Mouse FIL-1 theta	Mouse interleukin-	iCIL-1ra, Homo sa	ter	Human FIL-1 theta	Human interleukin-	Human interleukin-	Human IL-1 delta h
225	AAY45062	AAE06655	AAB87601	AAB35260	AAB35262	999	AAY43526	AAB35263	AAB35264	AAY92256	AAY92254	AAY92253	AAY92255	ABG22711	AAY97068	AAB35266	AAY43525	AAB66663	AAY97067	AAM50219	AAY33281	AAB66665	AAB84999	AAY43527	AAM50220	AAE05841	AAB19925	AAR73642	AAM50217	AAE05840	AAB19922	92	ABB11747
21	21	22	22	22	22	22	50	22	22	21	21	21	21	22	21	22	20	22	21	22	50	22	22	20	22	22	22	16	22	22	22	22	22
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ALIGNMENTS

RESULT AAW862	RESULT 1 AAW96284
Q X	AAW86284 standard; Protein; 156 AA.
X AC	AAW86284;
TO X	19-FEB-1999 (first entry)
DE X	Rodent interleukin (IL)-1 delta polypeptide.
K K	Interleukin; IL-1 delta; polyclonal antibody; IL-1 epsilon; cytokine;
KW.	chemokine.
SO	Mus sp.
NA S	WO9847921-A1.
PD CA	29-OCT-1998.
PF V	17-APR-1998; 98WO-US06879.
PR	06-AUG-1997; 97US-0055111.
R X	21-APR-1997; 97US-0837627.
PA	(SCHE) SCHERING CORP.
PI	Bazan JF, Hedrick JA, Kastelein RA, Sana TR;
DR X	WPI; 1998-609976/51. N-PSDB; AAV71958.
F F	Mammalian interleukin 1-delta and 1-epsilon - useful for, e.g.
7 3	reguranting cue immune system and inilammatory responses

Mammalian interleukin 1-delta and 1-epsilon - useful for, e.g. regulating the immune system and inflammatory responses

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Claim 1; Page 67; 72pp; English.
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                                      This represents a rodent interleukin (IL)-1 delta polypeptide. The invention relates to a recombinant polypeptide that specifically binds polyclonal antibodies (Abs) generated against a 12 consecutive amino acid segment of IL-1 delta or IL-1 epsilon. Agonists or antagonists of these IL polypeptides are used to regulate a cell involved in an inflammatory response. The IL-1 delta or IL-1 epsilon polypeptides and peptides and sused to produce Abs and antigen-Ab complexes. The polypeptides, Abs and the corresponding nucleic acids regulate development and/or the immune system, and can be used to diagnose and treat conditions associated with abnormal expression of IL. Agonists or antagonists of IL-1 delta or IL-1 epsilon polypeptides are used with agonists or antagonists of IL-1 alpha, IL-1 Bret, IL-1 gamma, IL-2 and/or IL-12. The IL-1 delta or IL-1 epsilon polypeptides may be used as a soluble polypeptide or as a fusion protein with another cytokine or chemokine.
                                                                                                                                                                                                                                                                                                                                                                                                                  61 GVOGGSOCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 120
                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                               1 MMVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVIL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Interleukin 1 delta; IL-1 delta; glaucoma; ectodermal dysplasia;
insulin-dependent diabetes mellitus; wrinkly skin syndrome;
T-cell leukemia; lymphoma; tibial muscular dystrophy.
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                                                                                                                                                                                                                                                                                        Query Match 100.0%; Score 819; DB 19; Length 156; Best Local Similarity 100.0%; Pred. No. 2.7e-87; Matches 156; Conservative 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         121 FLCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Pages 89-90; 113pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY28407 standard; Protein; 156 AA
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98US-0071074.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-458310/38.
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                                                                                                                                                                                                                                                      Sequence
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The present sequence represents mouse interleukin 1 delta (IL-1 delta).

IL-1 delta proteins are useful for the determination of the molecular weight of a sample protein. The protein and its fragments are useful as controls for peptide fragmentation. This is useful for determining the isoelectric point of a sample protein. Antibodies generated against IL-1 delta and its fragmented peptides can be used to enhance the accuracy of these molecular weight markers to determine the apparent molecular aight and isoelectric point of a sample protein. IL-1 delta conter-structure molecules. IL-1 delta associated with IL-1 delta counter-structure molecules. IL-1 delta can also be used as therapeutic agents for the treatment of diseases mediated by IL-1 delta IL-1 delta may be used as a reagent in studying the interleukin I delta coding sequences can be used to identify human chromosome 2, and to identify genes associated with certain diseases, especially with region 2dil-12, including glaucoma, ectodermal dysplasia, insulin-dependent diabetes mellitus, wrinkly skin syndrome, T-cell leukemia/lymphoma and tibial muscular dystrophy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Generic; interleukin-1; IL-1; homologue; zilla3; anti-inflammatory; antagonist; pro-inflammatory; agonist; immunomodulator; antiarthritic; antirheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic; immunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GVQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 MMVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVIL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 156;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.0%; Score 819; DB 20; Best Local Similarity 100.0%; Pred. No. 2.7e-87; Matches 156; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FLCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY92260 standard; Protein; 156 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Murine IL-1 homologue, zilla3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-US23533.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          West RR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-303780/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 156 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB; AAA09198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vulnerary; 2q14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-OCT-1998;
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This shows an interleukin-1 (IL-1) homologue, designated zilla3. A 350 bp probe generated from the DNA sequence by PCR using AAA09199-200 was used to analyze human northern blots.

It is believed that zilla3 acts through IL-1 receptors. In general, zilla3 proteins having a Lys residue at position 148 will have anti-inflammatory activity (e.g. AAY92256), whilst those having Asp (see AAY92254) or Glu at this position will have pro-inflammatory action. Zilla3 is used to modulate an immune response in an animal (claimed). Antagonists zilla3 forms may be used to treat or prevent chronic inflammatory diseases such as rheumatoid arthritis, of claimed). Antagonists zilla3 forms may be used to treat or prevent chronic inflammatory diseases such as rheumatoid arthritis, osteoarthritis and Lyme arthritis, psoriasis, to reduce tissue damage after ischemia, to treat septic shock, graft versus-host disease and leukemia. The antagonists may also alleviate inflammatory bowel disease including Crohn's disease and ulcerative colitis, insulin-dependent diabetes mellitus, acute pancreatitis, glomerulonephritis and cerebral ischemia. Agonist forms of zilla3 may promote wound healing by IL-1 effects on growth factor secretion and cell proliferation. They may also treat infections, especially gastrointestinal infections.
                              Proteins useful for treatment of inflammatory conditions such as rheumatoid arthritis and psoriasis are agonists or antagonists forms of
                                                                                                                                                    Example 7; Page 59-60; 64pp; English.
                                                                                              new interleukin-1 homologue
SSSSXS
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156 AA; Sequence

They may also

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                                                                                                     GVOGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 120
                                                                                                                    61 gvqgggqclscgtekgpilklepvnimelylgakesksftfyrrdmgltssfesaaypgw 120
                                Gaps
                                                    1 MMVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVIL 60
                                                                 ö
                               Indels
    Score 819; DB 21;
Pred. No. 2.7e-87;
0; Mismatches 0;
                                                                                                                                                      121 FLCTSPEADQPVRLTQIPEDRAWDAPITDFYFQQCD 156
                                                                                                                                                                    100.0%;
Query Match 100.
Best Local Similarity 100.
Matches 156; Conservative
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AAY45061; AY45061

AAY45061 standard; Protein; 156 AA.

(first entry) 31-MAY-2000

Murine TANGO-93 protein.

TANGO-93; cytokine; mouse; secreted protein; IL-1 expression; cancer; Interleukin-1 receptor antagonist; IL-1ra; inflammation; antiasthmatic; immunosuppressive; antirheumatic; antiarthritic; antiporiatic; asthma; antiinflammatory; antibacterial antiulicer; cytostatic; immunomodulator; osteopathic; dermatological; antidiabetic; psoriasis; ulcerative collitis; osteopathic; dermatological; antidiabetic; psoriasis; ulcerative collitis; septic shock; cachexia; Crohn's disease; chronic myelogenous leukaemia; septic shock; cachexia; crohn's disease; chronic myelogenous leukaemia; liver disease; diabetes; osteoarthritis; Hodgkin's disease; lyme disease; suctommune disease; myasthenia gravis; pharmacogenomic; diagnosis; systemic lupus erythematosus; forensic; transgenic animal.

WO200008045-A2.

17-FEB-2000

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The present sequence is the murine TANGO-93, a secreted protein, that belongs to the cytokine superfamily. It plays a role similar to secreted interleukin-1 receptor antagonist (IL-Ira) and ils expression is developmentally regulated in liver, heart and bone marrow. TANGO-93 modulates immune mediated inflammation and IL-1 gene or protein cypression. TANGO-93 is useful as a modulating agent for regulating cellular processes like asthma, graft vs-host disease, rheumatoid arthritis, psoriasis, inflammatory bowel disease, septic shock, ulcerative colitis, Croin's disease, chronic myelogenous leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer, liver disease, Hodgkin's disease, osteoarthritis, Lyme disease, cachexia, and autoimmune diseases e.g. myasthenia gravis, autoimmune diseases e.g. myasthenia gravis, autoimmune diabetes and systemic lupus erythematosus. Partial TANGO-93 sequences are useful in forensic biology, for diagnostic and prognostic assays, prophylactic and therapeutic treatment and pharmacogenomics. The DNA sequences are useful as hybridisation probes and primers, for isolation of TANGO-93 sequence and for the creation of transgenic animals.
                                                                                                                                                                                                                                                Isolated nucleic acid sequences encoding TANGO-93 polypeptide useful for treating a variety of cellular processes e.g. asthma, rheumatoid arthritis, psoriasis and autoimmune diseases
                                                                                    (MILL-) MILLENNIUM BIOTHERAPEUTICS INC.
                                                                                                                                                                                                                                                                                                                                      Claim 9; Fig 1; 113pp; English
99WO-US17886.
                                           98US-0131263
                                                                                                                                                                          WPI; 2000-205669/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               156 AA;
                                                                                                                                                                                                       N-PSDB; AAZ50811
                                      07-AUG-1998;
36-AUG-1999;
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                                                                                                                                    an X;
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ö Gaps 1 MMVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVIL 60 ; 0 Length 156; Indels Score 819; DB 21; Pred. No. 2.7e-87; Mismatches 0; ; 100.0%; 100.08; Conservative Query Match Best Local Similarity Matches 156; Conserv ð

61 GVQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRDMGLTSSFESAAYPGW 120 9 g ò

121 FLCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 156 ð g

g

AAY96937 standard; Protein; 155 S AAY96937 RESULT

AA.

(first entry)

31-OCT-2000

AAY96937;

Murine IL-1 receptor antagonist 3.

mIL-1Ra3; interleukin-1 receptor antagonist-3; IL-1lp; osteopathic; interleukin-1-like polypeptide; anti-inflammatory; anti-asthmatic; anti-arthritic; antimicrobial; respiratory; anti-ischemic; vaccine; dermatalogical; immunomodulatory; gastrointestinal; gene therapy.

WO200039297-A2.

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The present invention provides the protein and coding sequences of the human and murine interleukin-ILI (IL-ILI) proteins. The IL-ILI gene is located between the IL-Ibeta and IL-Ireceptor genes at human chromosome 2q13. The sequences are useful in the diagnosis, prevention and treatment of heart disease, cancer and inflammatory diseases such as rheumatoid arthritis, systemic lupus exythematosus, inflammatory bowel disorder, diabetes, psoriasis, osteoporosis, lichen sclerosis, ulcerative colitis, severe periodontal disease and pregnancy complications. The present sequence is the murine IL-ILI protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             62 VQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human and murine interleukin-1L1 polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 MYLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99.4%; Score 814; DB 22; Length 155; 100.0%; Pred. No. 1e-86; 1ve 0; Mismatches 0; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          interleukin; IL-1 receptor; cancer; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammatory processes
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                                                                                                                                                                                                                                                                                                                                                              (INTE-) INTERLEUKIN GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 11; Fig 3; 150pp; English
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                                                                                                                                                                                                                            17-JUL-2000; 2000WO-US19508
                                                                                                                                                                                                                                                                                               99US-0144298
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acids encoding useful for controlling
                                                                                                                                                                                                                                                                                                                                                                                                                             Nicklin M, Barton J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-091974/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 155; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  155 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200102571-A2.
                                                                                              WO200105974-A2.
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                                                                                                                                                                                                                                                                                               16-JUL-1999;
                                                                                                                                                             25-JAN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      An isolated nucleic acid molecule encoding an interleukin-1-like polypeptide (IL-11p) that retains one or more activities of the peptide from which it is derived, such as the IL-18R binding activity of a human interleukin-1 receptor antagonist-1 (hIL-1Ral) polypeptide, is new. The nucleic acids may be used in molecular engineering applications, e.g. hybridization assays and chromosome and gene mapphing studies, for recombinantly producing the IL-11p polypeptide or for producing gene knock out animals to study the role of the protein in metabolism and disease processes (conversely, gene therapy protocols may be used to supplement a patients production of the polypeptide or to rectify mutations that lead to the production of in active peptides). The peptides produced may be used to screen for and produce modulators (e.g. antibodies) of IL-11p protein expression and activity, butch may be use to treat disorders associated with inappropriate IL-11p expression and activity, such as inflammatory disorders, asthma, arthritis, sepsis, acute lung injury, adult respiratory distress standome, idiopathic pulmonary fibrosis, isohemic reperfusion disease. Soriansis, graft versus host disease and/or inflammatory bowel disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 vqgqsqclscgtekgpilklepvnimelylgakesksftfyrrdmgltssfesaaypgwf 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acids encoding interleukin-1-like polypeptides, useful for preventing and treating e.g. inflammation, asthma and psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    122 LCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 22; Fig 9A-B; 143pp; English.
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                                                                                                                              98US-0113430.
99US-0116843.
99US-0129122.
                                                               99WO-US30720.
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Best Local Similarity 100.
Matches 155; Conservative
                                                                                                                                                                                                                                                                   (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-452395/39
                                                                                                                                                                                                                                                                                                                                     Goddard A, Pan J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          155 AA;
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                                                               22-DEC-1999;
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22-JAN-1999;
                                                                                                                                                                                                13-APR-1999;
06-JUL-2000
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AAB35261
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The invention relates to murine interleukin-1 homologue 3 (IL-1H3; AB48B28) and nucleic acids which encode it (CDNA given in AAC81700), including nucleic acids equences with at least 95% identity to AAC81700. The invention also relates to expression vectors and host cells comprising nurine IL-1H3 nucleic acids, the recombinant production of murine IL-1H3 agonists and antagonists thus identified. IL-1H3 agonists and antagonists are of use for treating human diseases such as chronic or inflammatory bowel disease, psorians and arthritis), transplant rejection, graft versus host disease, infection, stroke, ischaemia, cute respiratory distress syndrome, allergies, asthma, restenosis, brain injury, AIDS, bone diseases (e.g., osteoporosis), cancers atherosclerosis, and Alzheimer's disease. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse; interleukin-idelta; IL-idelta; virucide; hepatotropic; fever; immunological disorder; tumour; inflammatory disorder; hypoglycaemia; autoimmune disease; pulmonary tuberculosis; fulminant hepatitis; leprosy; psoriasis; viral infection; allergy; cytokine; HIV; drug screening.
                                                                                                                                                                                                                                                      Murine interleukin-1 homolog polypeptide used for screening modulators of the polypeptide which can be used for treating autoimmune diseases, cancer, brain injury and bone disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 62 VQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22; Length 155;
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100.0%; Pred. No. 1e-86;
ive 0; Mismatches
                                                                                                                                                Halsey W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse interleukin-1delta (IL-1delta) protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             122 LCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      McDonnell PC,
                                                                                                                                                                                                                                                                                                                                             Claim 2; Page 28-29; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAE06663 standard; Protein; 154 AA.
                                                                           (SMIK ) SMITHKLINE BEECHAM CORP. (SMIK ) SMITHKLINE BEECHAM PLC.
         99US-0135599.
2000US-0577715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 99.4
Best Local Similarity 100.
Matches 155; Conservative
                                                                                                                                              Smith RF, Young PR,
                                                                                                                                                                                       WPI; 2001-025138/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               155 AA;
                                                                                                                                                                                                              N-PSDB; AAC81700
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             24-MAY-1999;
23-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Interleukin-1 homologue 3; IL-1H3; mouse; murine; drug screening; agonist; antagonist; human disease; chronic inflammation; acute inflammation; septicaemia; autoimmune disease; psoriasis; inflammatory bowel disease; arthritis; transplant rejection; infection; graft versus host disease; arthritis; transplant rejection; infection; graft versus host disease; acute respiratory distress syndrome; allergy; asthma; restenosis; stroke; ischaemia; brain injury; AIDS; bone disease; osteoporosis; cancer; lymphoproliferative disorder; atherosclerosis; congestive heart failure; Aizheimer's disease; immunosuppressive;
                                                                                                                                                                                                                                        Isolated nucleic acids encoding interleukin-1 (IL-1) receptor antagonist proteins (referred as IL-1Hy1), useful in the treatment of cancer, e.g. breast adenocarcinoma and brain tumors, and an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      62 VQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61
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                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to interleukin (IL)-1 receptor antagonist proteins. IL-1Hy1 is useful for treating cancer, an inflammatory disease mediated by IL-18, inflammation resulting from infection or allergic reactions, and inflammation associated with chronic bronchitis, arthritis, diabetes or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 814; DB 22; Length 155;
Pred. No. 1e-86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 156
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murine interleukin-1 homologue 3 (IL-1H3).
                                                                                                                                                                                                                                                                                                             inflammatory disease mediated by IL-18
                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 177; 179pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99.4%; Scc.
100.0%; Pre/
0; }
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antimicrobial; neuroprotective.
13-OCT-1999; 99US-0417455.
08-DEC-1999; 99US-0457626.
10-MAR-2000; 2000US-0523552.
22-MAY-2000; 2000US-0576008.
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                                                                                                                                                                                              WPI; 2001-071582/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     155 AA;
                                                                                                              (HYSE-) HYSEQ INC.
                                                                                                                                                       Pace A;
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Query Match

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Best Local

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AAB48828 RESULT

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hIL-1Ra3; human interleukin-1 receptor antagonist-3; IL-11p; osteopathic;
                                                                                                           98US-0087393
                                                                              99WO-US00514
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Best Local Similarity 91.0 Matches 141; Conservative
                                                                                                                                                 (IMMV ) IMMUNEX CORP.
                                                                                                                                                                                                    WPI; 1999-458310/38.
N-PSDB; AAX89432.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         155 AA;
 Homo sapiens.
                           WO9935268-A1
                                                                                                         01-JUN-1998;
09-JAN-1998;
                                                                               08-JAN-1999;
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                                                    15-JUL-1999
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                                                                                                                                                                                                                                                                   molecules and their corresponding nucleic acid sequences, designated as interleukin-idelta (Ir-idelta) and interleukin-lepsilon (Ir-lepsilon).

IL-idelta and II-lepsilon are useful for treating conditions exhibiting abnormal expression of the interleukin such as immunological disorders, tumours, inflammatory disorders, fever, hypoglycaemia, psoriasis, allery, autofinance diseases and infectious diseases (e.g., pulmonary tuberculosis, leprosy, fulminant hepatitis, and viral infections such as HIV). The invention also relates to methods of using the composition containing II-ldelta or IL-lepsilon for both diagnostic and therapeutic utilities. IL-ldelta or IL-lepsilon for both diagnostic and therapeutic utilities. IL-ldelta specific, e.g., capable of distinguishing between IL-l family members and an IL-idelta, for the interleukin or its antibodies specific and insurance of elevated levels of expression, or immunological disorders which lead to antibody production contemplates or immunological disorders which lead to antibody production competitive drug screening assays. The presence of elevated levels of competitive drug screening assays. The present sequence is mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                              Novel isolated or recombinant antigenic interleukin-1 delta or epsilon polypeptide useful for treating conditions exhibiting abnormal expression of interleukin such as immunological disorders, tumor and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    63 QGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFL 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                            The invention relates to recombinant antigenic interleukin-1 like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Interleukin 1 delta; IL-1 delta; glaucoma; ectodermal dysplasia;
insulin-dependent diabetes mellitus; wrinkly skin syndrome;
T-cell leukemia; lymphoma; tibial muscular dystrophy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 154;
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                                                                                                            Kastelein RA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98.8%; Score 809; DB 22;
100.0%; Pred. No. 3.8e-86;
iive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bazan JF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY28408 standard; Protein; 155 AA
                                                                                                                                                                                                                              Disclosure; Fig 1; 103pp; English.
                         01-FEB-2001; 2001WO-US03285.
                                                    02-FEB-2000; 2000US-0179638.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human interleukin 1 delta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches 154; Conservative
                                                                                                           Debets JEMA, Timans JC,
                                                                               SCHE ) SCHERING CORP
                                                                                                                                      WPI; 2001-488886/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
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09-AUG-2001
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                                                                                                                                                                                                                                                       Weight of a sample protein. The protein and its fragments are useful controls for peptide fragmentation. This is useful determining the controls for peptide fragmentation. This is useful for determining the isoelectric point of a sample protein. This is useful for determining the isoelectric point of a sample protein. Antibodies generated against IL-1 delta and its fragmented peptides can be used to enhance the accuracy of these molecular weight markers to determine the apparent molecular eight and isoelectric point of a sample protein. IL-1 delta can be used to screen for potential inhibitors of activity associated with IL-1 delta counter-structure molecules. IL-1 delta can also be used as therapeutic agents for the treatment of diseases mediated by IL-1 delta delta may be used as a reagent in studying the interleukin it. It.1 delta counter-structure molecules. IL-1 signalling pathway, or as a reagent to block IL-1 signalling. The interleukin it is delta coding sequences can be used to identify human chromosome 2, and to identify name chromsome 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and to identify genes associated with certain diseases, especially with region 2q11-12, including glaucoma, ectodermal dysplasia, insulindependent diabetes mellitus, wrinkly skin syndrome, T-cell leukemia/lymphoma and tibial muscular dystrophy.
                                                                                                                                                                                                                               The present sequence represents human interleukin 1 delta (IL-1 delta).
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Murine and Human interleukin 1 delta DNA, polypeptides and 1ts fragments, useful as molecular weight markers
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antagonist; pro-inflammatory; agonisť; immunomodulator; antiarthritic; antirheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic; immunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
                                                                    Seneric; interleukin-1; IL-1; homologue; zilla3; anti-inflammatory;
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131..133
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/label= loop
66..71
/label= beta-strand
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/label- beta-strand
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label= beta-strand
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label= beta-strand
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/label- beta-strand
                                                                                                                                                                                         Location/Qualifiers
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/label- beta-strand
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/label= beta-strand
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/label- beta-strand
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/label= beta-strand
                                      Human IL-1 homologue, hzilla3.
                                                                                                                                                                                                                                    14..16
/label- loop
                                                                                                                                                                                                                                                                                            2..25
label= loop
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/label= loop
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/label- loop
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/label= loop
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/label 100p
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/label- loop
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/label= loop
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         10-AUG-2000 (first entry)
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                                                                                                                                                          Homo sapiens.
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       An isolated nucleic acid molecule encoding an interleukin-1-like polypeptide (IL-11p) that retains one or more activities of the peptide from which it is derived, such as the IL-18R binding activity of a human interleukin-1 receptor antaqonist-1 (hIL-1Ral) polypeptide, is new. The nucleic acids may be used in molecular engineering applications, e.g. hybridization assays and chromosome and gene mapping studies, for recombinantly producing the IL-11p polypeptide or for producing gene concesses (conversely, gene therapy protocols may be used to supplement a patients production of the polypeptide or to rectify mutations that lead to the production of the polypeptides). The peptides produced may be used to supplement any be used to screen for and produce modulators (e.g. antibodies) of IL-11p protein expression and activity which may be use to treat disorders associated with inappropriate IL-11p expression and activity which may be use activity, we associated with inappropriate IL-11p expression and activity, which may be use such as inflammancy disorders, asthritis, osteoarthritis, sepsis, acute lung injury, adult respiratory distress syndrome, idiopathic pulmonary fibrosis, ischemic reperfusion disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
interleukin-1-like polypeptide; anti-inflammatory; anti-asthmatic;
anti-arthritic; antimicrobial; respiratory; anti-ischemic; vaccine;
dermatalogical; immunomodulatory; gastrointestinal; gene therapy.
                                                                                                                                                                                                                                                                                                                                             Nucleic acids encoding interleukin-1-like polypeptides, useful for preventing and treating e.g. inflammation, asthma and psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 734; DB 21; Length 155;
Pred. No. 2.1e-77;
4; Mismatches 10; Indels (
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                                                                                                                                                                                                                                                                                                                                                                                        Claim 22; Fig 7; 143pp; English.
                                                                                                                                                                            98US-0113430.
99US-0116843.
99US-0129122.
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91.0%;
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Best Local Similarity 91.0°
Matches 141; Conservative
                                                                                                                                                                                                                                         (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                               WPI; 2000-452395/39.
N-PSDB; AAA51597.
                                                                                                                                                                                                                                                                       Goddard A, Pan J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         155 AA;
                                                                                       WO200039297-A2.
                                                              Homo sapiens.
                                                                                                                                                   22-DEC-1999;
                                                                                                                                                                               23-DEC-1998;
                                                                                                                                                                                                             13-APR-1999;
                                                                                                                      06-JUL-2000
                                                                                                                                                                                               2-JAN-1999;
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AAY92257 RESULT

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Human; interleukin-1delta; IL-1delta; virucide; hepatotropic; fever;
           Homo sapiens.
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                                                                                                                                                                                                               This shows an interleukin-1 (IL-1) homologue, designated zilla3.

It is believed that zilla3 acts through IL-1 receptors. In general, zilla3 perclains waving a Lys residue at position 148 will have anti-inflammatory activity (e.g. AAY92256), whilst those having Asp (see AAY92254) or Glu at this position will have pro-inflammatory action. Zilla3 is used to modulate an immune response in an animal claimed). Antagonists zilla3 forms may be used to treat or prevent chronic inflammatory diseases such as rheumatorid arthritis, osteoarthritis and Lyme arthritis, psoriasis, to reduce tissue damage after ischemia, to treat septic shock, graft-versus-host disease and including Crohn's disease and ulcerative colitis, insulin-dependent diabetes mellitus, acute pancreatitis, glomerulonephritis and cerebral ischemia. Aponist forms of zilla3 may promote wound healing by IL-1 treat infections, especially gastrointestinal infections.
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                                                                                                                                                                                                                                                                                                                                                                                                                   They may also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TANGO-93; cytokine; human; secreted protein; IL-1 expression; cancer; Interleukin-1 receptor antagonist; IL-1ra; inflammation; antiasthmatic; immunosuppressive; antirheumatic; antiarthritic; antiporiatic; asthma; antiinflammatory; antibacterial; antiulcer; cytostatic; immunomodulator; osteopathic; dermatological; antidiabetic; psoriasis; ulcerative colitis; graft vs-host disease; rheumatoid arthritis; inflammatory bowel disease; septic shock; cachexia; Crohn's disease; chronic myelogenous leukaemia; liver disease; diabetes; osteoarthritis; hodgkin's disease; Lyme disease; cutoimmune disease; myasthenia gravis; pharmacogenomic; diagnosis; chromosome 2; systemic lupus erythematosus; forensic; transgenic animal.
                                                                                                                                   Proteins useful for treatment of inflammatory conditions such as rheumatoid arthritis and psoriasis are agonists or antagonists forms of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 155;
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Pred. No. 2.1e-77;
4; Mismatches 10;
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                                                                                                                                                                                           Disclosure; Page 52-53; 64pp; English.
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                                                                   Clegg CH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    89.6%;
91.0%;
           98US-0169745
                                                                                                                                                                new interleukin-1 homologue
                                                                                                          N-PSDB; AAA09193, AAA09194
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Best Local Similarity 91.0
Matches 141; Conservative
                                       (ZYMO ) ZYMOGENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human TANGO-93 protein.
                                                                   West RR,
                                                                                            2000-303780/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                            155 AA;
           08-OCT-1998;
                                                                   Sheppard PO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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The present sequence is the human TANGO-93, a secreted protein that
belongs to the cytokine superfamily. It plays a role similar to secreted
conterlenkin-1 receptor antagonist (IL-Ira) and its expression is
conterlenkin-1 receptor antagonist (IL-Ira) and its expression is
developmentally regulated in the uterus, placenta and skeletal muscles.
Human TANGO-93 gene is mapped to chromosome 2 within the IL-I cluster.
TANGO-93 modulates immune mediated inflammation and IL-1 gene or protein
expression. TANGO-93 is useful as a modulating agent for regulating
callular processes like asthma, graft vs-host disease, rheumatoid
arthritis, psoriasis, inflammatory bowel disease, septic shock,
ulcerative colitis, Crohn's disease, chronic myelogenous leukaemia,
cacheria, and autoimmune diseases, chronic myelogenous leukaemia,
cacheria, and autoimmune diseases e.g. myasthenia gravis, autoimmune
diabetes and systemic lupus erythematosus. Partial TANGO-93 sequences
are useful in forensic biology, for diagnostic and prognostic assays,
sprophylactic and therapeutic treatment and pharmacogenomics. The DNA
sequences are useful as hybridisation probes and primers, for isolation
control of TANGO-93 sequence and for the creation of transgenic animals.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Isolated nucleic acid sequences encoding TANGO-93 polypeptide useful for treating a variety of cellular processes e.g. asthma, rheumatoid arthritis, psoriasis and autoimmune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Pred. No. 2.1e-77;
4; Mismatches 10; Indels
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                                                                                                                                                                                                                                                                           (MILL-) MILLENNIUM BIOTHERAPEUTICS INC
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91.0%;
                                                                                                                                     99WO-US17886.
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Best Local Similarity 91.0
Matches 141; Conservative
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WO200008045-A2
                                                                                                                                                                                                          07-AUG-1998;
                                                                                                                                         06-AUG-1999;
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                                                                    17-FEB-2000
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immunological disorder; tumour; inflammatory disorder; hypoglycaemia;
autoimmune disease; pulmonary tuberculosis; fulminant hepatitis; leprosy;
psoriasis; viral infection; allergy; cytokine; HIV; drug screening.
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/note= "IL-1 receptor beta subunit binding region"
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te= "IL-1 receptor beta subunit binding region"
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re= "Beta sheet region"
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26..29
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The invention relates to recombinant antigenic interleukin-1 like molecules and their corresponding nucleic acid sequences, designated as interleukin-delata and interleukin-lepsilon (IL-lepsilon).

IL-idelta and IL-lepsilon are useful for treating conditions exhibiting abnormal expression of the interleukin such as immunological disorders, tumours, inflammatory disorders, fever, hypoglycaemia, psoriasis, allergy, autoimmune diseases and infectious diseases (e.g., pulmonary tuberculosis, leprosy, fulminant hepatitis, and viral infections such as HIV). The invention also relates to methods of using the composition containing IL-ldelta or IL-lepsilon for both diagnostic and therapeutic utilities. IL-ldelta for IL-lepsilon for both diagnostic and therapeutic utilities. IL-ldelta is used as an immunogen for the production of antisera or antibodies specific, e.g., capable of distinguishing between IL-ldelta, for the interleukin or its ragment. The purified interleukin is used as a reagent to detect any antibodies generated in response to the presence of elevated levels of expression, or immunological disorders which lead to antibody production competitive drug screening assays. The present sequence is human interleukin-ldelta (IL-ldelta) protein.
                                                                                                                                                                                                                                                                                                                                                 Novel isolated or recombinant antigenic interleukin-1 delta or epsilon polypeptide useful for treating conditions exhibiting abnormal expression of interleukin such as immunological disorders, tumor and
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         /note= "IL-1 receptor beta subunit binding region"
155
                                                 /note= "IL-1 receptor beta subunit binding region"
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Pred. No. 2.1e-77;
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/note= "IL-1 receptor alpha subunit binding region"

Job time: 210 sec

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eighty four nucleic acids encoding PRO polypeptides, useful in molecular biology, including use as hybridization probes, and in chromosome and gene mapping.
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Wood WI;
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Watanabe CK,
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99WO-US21090.
99US-0169495.
99US-0170262.
2000US-0175481.
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2000WO-US04414.
2000WO-US05601.
                                                   Human; PRO protein; mapping
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J, Gurney AL,
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N-PSDB; AAF92133.
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Best Local Similarity
Matches 141; Conserv
                                                                                                     WO200116318-A2.
                          Human PRO4342
                                                                            Homo sapiens.
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Grimaldi CJ,
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15-SEP-1999;
07-DEC-1999;
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05-JUN-2000;
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US-09-770-528-2 819 1 MMVLSGALCFRMKDSALKVL.....IPEDPAWDAPITDFYFQQCD 156 Title: Perfect score: Sequence:

Scoring table:

231628 seqs, 24425594 residues BLOSUM62 Gapop 10.0 , Gapext 0.5 Searched:

231628 tal number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

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Listing first 45 summaries

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		Appli	Appli	App1i	Appli	Appli	Appli	Appli	, Appl		Appli	Appli	Appli	App11	Appli	, Appl	, Appl	, Appl	Appli	Appli	Appli	Appli	Appli	Appli	Appli	Appli		, Appl
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SUMMARIES	ΩI	US-09-417-455	US-09-348-942-5	US-09-31	US-09-417-455-3	US-09-348-942-3	US-09-417-455-9	US-09-348	DS-000-60-SD	US-08-862-730C	US-09-31	US-09-316-081-	US-08-798-414-	US-09-131-247-2	US-09-131-247-4	US-08-476	US-08-91	US-08-910-884-10	US-08-459-811-2	US-08-484	US-08-459-092-2	US-08-459-814-2	US-08-425-232-	US-08-471-227	US-08-479	US-08-477-143-	US-09-417-455-1	US-09-348
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Sequence 9, Appli Sequence 2, Appli Sequence 2, Appli Sequence 20, Appl Sequence 10, Appl Sequence 11, Appl Sequence 13, Appl Sequence 13, Appl Sequence 13, Appl Sequence 13, Appl Sequence 14, Appl Sequence 16, Appl Sequence 16, Appl Sequence 16, Appl Sequence 16, Appl Sequence 23, Appl Sequence 23, Appl	SRECEPTOR ANTAGONIST AND USES THEREOF	Length 155; ; Indels 0; Gaps 0; SVVPNRALDASLSPVILG 61
4 US-09-316-081-9 1 US-08-42-655-2 2 US-08-080-185-2 3 US-09-000-630C-20 3 US-09-000-630C-20 3 US-09-128-155-14 4 US-09-141-455-30 4 US-09-316-081-8 4 US-09-316-081-8 4 US-09-316-081-8 4 US-09-316-081-8 4 US-09-316-081-8 5 US-08-910-733-13 5 US-08-910-733-13 7 US-08-910-733-13 7 US-08-910-733-13 7 US-08-910-733-13 7 US-08-910-733-13 7 US-09-131-247-14	MENT 1.N-1 2.2 2.2 2.2 2.2 2.0 0.0 0.0 0.0 0.0 0.0	%; Score 734; DB 4; 9%; Pred. No. 8.1e-86; 4; Mismatches 10 LHNNQLLAGGLHAEKVIKGEEI
37.5 37.5 37.5 37.5 37.5 177 37.5 177 37.5 177 177 177 177 177 177 177 177 177 17	5 Application US/294655 RMATION: FORD JOHN VENTION: FORD JOHN VENTION: A NOVE INCE: 28110/363 LICATION NUMBER: G DATE: 1999-04 CATION NUMBER: G DATE: 1999-04 CATION NUMBER: G DATE: 1999-04 CATION NUMBER: G DATE: 1999-07 CATION NUMBER: G DATE: 1999-07 CATION NUMBER: G DATE: 1998-05 CATION NUMBER: G DATE: 1998-	89.61 Similarity 91.03 41; Conservative T.SGALCFRMKDSALKVLXI LLSGALCFRMKDSALKVLXI GGSQCLSCGTEKGPILKLI
228 229 330 330 330 330 330 330 330 33	RESULT 1 US-09-417-455- Sequence 5, Patent No. 6 GENERAL INFO APPLICANT: APPLICANT: TITLE OF IN TITLE OF IN TITLE OF IN FILE REFERE CURRENT APPLI PRIOR FILIN PRIO	Query Match Best Local S Matches 141 Oy 2 WULS Oy 1 Db 1 MVLS OY 62 VOG

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62 VQGGSQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 121
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91.0%; Pred. No. 8.1e-86;
iive 4; Mismatches 10
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SEQ ID NO 5
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Matches 141; Conservative
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Best Local Similarity 90.07
Matches 72; Conservative
                                                                                                ; ORGANISM: Homo sapiens
US-09-316-081-5
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APPLICANT: Ford, John
APPLICANT: Pace, Ann
                                                       LENGTH: 155
TYPE: PRT
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TITLE OF INVENTION: A NOVEL INTERLEUKIN-1 RECEPTOR ANTAGONIST AND USES THEREOF
FILE REFERENCE: 28110/35801
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61 VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
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Patent No. 6339141
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ballinger, Dennis G.
APPLICANT: Pace, Ann M.
TITLE OF INVENTION: Interleukin-1 Hy2 Materials and Methods
FILE REFERENCE: 28110/35659
CURRENT APPLICATION NUMBER: US/09/316,081
CURRENT FILING DATE: 1999-05-20
NUMBER OF SEQ ID NOS: 11
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Pred. No. 8.1e-86;
                                              122 LCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 156
                                                                     122 LCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 156
                                                                                                                                                                                                                                                                                                                        CURRENT PELLICATION NUMBER: US/09/348,942
CURRENT FILING DATE: 1999-07-07
EARLIER APPLICATION NUMBER: PCT/US99/04291
EARLIER PILING DATE: 1999-04-05
EARLIER FILING DATE: 1999-04-05
EARLIER PILING DATE: 1999-04-05
EARLIER FILING DATE: 1999-02-17
EARLIER APPLICATION NUMBER: US 09/251,370
EARLIER PILING DATE: 1999-02-17
EARLIER PILING DATE: 1999-07-31
EARLIER PILING DATE: 1999-07-31
EARLIER PILING DATE: 1998-07-31
EARLIER FILING DATE: 1998-05-10
EARLIER FILING DATE: 1998-04-03
NUMBER: OF SED ID NOSE: 10
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Patent No. 6337072
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Best Local Similarity 91.0%;
Matches 141; Conservative 4
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US-09-348-942-5
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APPLICANT: Pace, Ann
TITLE OF INVENTION: A MOVEL INTERLEUKIN-1 RECEPTOR ANTAGONIST AND USES THEREOF
FILE REFERENCE: 28110/36328
CURRENT APPLICATION NUMBER: US/09/417,455
CURRENT PELLING DATE: 1999-10-13
PRIOR PELLING DATE: 1999-04-05
PRIOR FILING DATE: 1999-04-05
PRIOR PELLING DATE: 1999-04-05
PRIOR FILING DATE: 1999-04-05
PRIOR PELLING DATE: 1999-04-13
PRIOR PELLING DATE: 1999-01-13
PRIOR PELLING DATE: 1999-05-19
PRIOR PELLING DATE: 1999-05-19
PRIOR PELLING DATE: 1999-05-10
PRIOR PELLING DA
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Pred. No. 1e-42;
3; Mismatches 5; Indels
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; ORGANISM: Mus musculus
US-09-417-455-9
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ORGANISM: Mus musculus
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LENGTH: 178
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Patent No. 6294655

GENERAL INFORMATION:
APPLICANT: FOLG. John
APPLICANT: FOLG. Ann
TITLE OF INVERTION: A NOVEL INTERLEUKIN-1 RECEPTOR ANTAGONIST AND USES THEREOF
FILE REPERENCE: 28110/36328
CURRENT APPLICATION NUMBER: US/09/417,455
CURRENT FILING DATE: 1999-10-13
PRIOR APPLICATION NUMBER: PCT/US99/04291
PRIOR APPLICATION NUMBER: PCT/US99/04291
PRIOR FILING DATE: 1999-04-05
                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: John Ford
TITLE OF INVENTION: A NOVEL INTERLEUKIN-1 RECEPTOR ANTAGONIST AND USES THEREOF
FILE REFERENCE: 28110/35801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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Pred. No. 1e-42;
3; Mismatches
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CURRENT PAPLICATION NUMBER: US/09/348,942

CURRENT FILING DATE: 1999-07-07

EARLIER APPLICATION NUMBER: US/09/291

EARLIER FILING DATE: 1999-04-05

EARLIER FILING DATE: 1999-04-05

EARLIER FILING DATE: 1999-04-05

EARLIER FILING DATE: 1999-04-05

EARLIER FILING DATE: 1999-01-13

EARLIER FILING DATE: 1998-05-19

EARLIER FILING DATE: 1998-05-19

EARLIER FILING DATE: 1998-05-10

EARLIER FILING DATE: 1998-05-15

EARLIER FILING DATE: 1998-05-10

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Patent No. 6337072
137 IPEDPAWDAPITDFYFQQCD 156
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90.0%;
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Best Local Similarity 90.0
Matches 72; Conservative
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US-09-348-942-3
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US-09-417-455-9
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APPLICANT: John Ford

TITLE OF INVENTION: A NOVEL INTERLEUKIN-1 RECEPTOR ANTAGONIST AND USES THEREOF
FILE REFERENCE: 28110/J580,
CURRENT APPLICATION NUMBER: US/09/348,942
CURRENT PILING DATE: 1999-07-07
EARLIER PILING DATE: 1999-07-07
EARLIER FILING DATE: 1999-04-05
EARLIER FILING DATE: 1999-04-05
EARLIER FILING DATE: 1999-04-05
EARLIER FILING DATE: 1999-02-17
EARLIER FILING DATE: 1999-02-17
EARLIER FILING DATE: 1999-01-13
EARLIER FILING DATE: 1999-07-01
EARLIER FILING DATE: 1999-07-01
EARLIER FILING DATE: 1998-07-31
EARLIER FILING DATE: 1998-07-31
EARLIER FILING DATE: 1998-07-31
EARLIER FILING DATE: 1998-06-19
EARLIER FILING DATE: 1998-06-19
EARLIER FILING DATE: 1998-06-19
EARLIER FILING DATE: 1998-06-15
EARLIER FILING DATE: 1998-04-03
NUMBER: OF SEO TO NOS : 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            39.0%; Score 319.5; DB 48.6%; Pred. No. 8.4e-33; ive 17; Mismatches 49
PRIOR FILLING DATE: 1999-01-13
PRIOR APPLICATION NUMBER: US 09/229,591
PRIOR PILING DATE: 1999-01-13
PRIOR APPLICATION NUMBER: US 09/127,698
PRIOR FILING DATE: 1998-07-31
PRIOR PLING DATE: 1998-07-31
PRIOR PELING DATE: 1998-06-19
PRIOR PELING DATE: 1998-06-19
PRIOR PILING DATE: 1998-05-10
PRIOR APPLICATION NUMBER: US 09/079,909
PRIOR APPLICATION NUMBER: US 09/079,909
PRIOR PILING DATE: 1998-05-15
PRIOR PELING DATE: 1998-05-15
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 9
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154 DRPVSLTNTPEEP---LIVTKFYFQE 176
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Patent No. 6337072
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Best Local Similarity 48.68
Matches 71; Conservative
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COUNTRY:
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US-09-316-081-2
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                                                                                 Gaps
                                                                                                                10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGVQGGSQCL 69
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                                                                                                                                                                                                                                                                                                                                                       Sequence 21, Application US/09000630C
Patent No. 6018029
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Fuentes, Nelson L.
TITLE OF INVENTION: DNA Encoding Canine Interleukin-1 Receptor TITLE OF INVENTION: Antagonist NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
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                                                DB 4; Length 178;
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                                                                               Indels
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                                                                               49;
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47.9%; Pred. No. 6.6e-32;
tive 17; Mismatches 50;
                                              39.0%; Score 319.5; DB 4
48.6%; Pred. No. 8.4e-33;
Live 17; Mismatches 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM compatible
OPERATING SYSTEM: Microsoft Windows
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE: mouse IL-1ra sequence US-09-000-630C-21
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/862,730
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: WordPerfect 6.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 21: SEQUENCE CHARACTERISTICS: LENGTH: 178 amino acids TYPE: amino acid STRANDEDNESS: single
                                            Query Match
Best Local Similarity 48.64
Matches 71; Conservative
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Best Local Similarity 47.9%
Matches 70; Conservative
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COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
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                                                                                                                                                                                                                                                                                                                                               US-09-000-630C-21
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US-09-348-942-9
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70 SCGTEKGPI-LKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTSPEA 128
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      39 FRIWDINOKIFYLRNNQLIAGYLQGPNIKLEGKIDMVP----IDLHSVFLGIHGGKLCL 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGVQGGSQCL 69
                                                                                                                    Nelson L.
DNA Encoding Canine Interleukin-1 Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6
                                                                                                                                                                                                              ADDRESSEE: Douglas C Murdock/ Bradley, Arant, Rose & White STREET: 2001 Park Place, Sulte 1400 CITY: Birmingham STATE: Alabama
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 178;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Ballinger, Dennis G.
APPLICANT: Ballinger, Dennis G.
APPLICANT: Pace, Ann M.
TITLE OF INVENTION: Interleukin-1 Hy2 Materials and Methods
FILE REFERENCE: 28110/35659
CURRENT APPLICATION NUMBER: US/09/316,081
CURRENT FILING DATE: 1999-05-20
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 38.2%; Score 312.5; DB 3; Best Local Similarity 47.9%; Pred. No. 6.6e-32; Matches 70; Conservative 17; Mismatches 50;
                                                                                                                                                                                                                                                                                                                                             ZIP: 3223-2736
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch,
COMPUTER: IBM compatible
OPERATING SYSTEM: Microsoft Windows
SOFTWARE: WordPerfect 6.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: mouse IL-1ra sequence US-08-862-730C-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US/08/862,730C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  129 DOPVRLTQIPEDPAWDAPITDFYFQQ 154
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US-08-862-730C-21; Sequence 21, Application US/08862730C
Faction to . 6063600; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Application US/09316081 Patent No. 6339141
                                                                                                                                                                   Antagonist
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/6
FILING DATE: 5/23/97
INFORMATION FOR SEQ ID NO: 21:
                                                                                       Gerald M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 178 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                       APPLICANT: Fuller, Gera APPLICANT: Fuentes, Nel: TITLE OF INVENTION: DNA TITLE OF INVENTION: Ant: NUMBER OF SEQUENCES: 27 CORRESPONDENCE ADDRESS:
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; ORGANISM; Human
US-09-131-247-2
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                                                                                                                                                                                                                                                                                                                                                               61 GVQGGSQCLSC-GTEKGPILKLEPVNIMELYLGAKESKSFTFYRDMGLTSSFESAAYPG 119
                                                                                                                                                                                                                                                                                                                                                                                                                    61 GVQGGSQCLSC-GTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPG 119
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                                                                                                                                                                               1 MMVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVIL 60
                                                                                                                                                                                                                                                   1 MCSLPMARYXIIKYADQKALYTRDGQLLVGDPVADNCC-AEKICTLPNRGLDRTKVPIFL 59
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         Length 152;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Ballinger, Dennis G.
APPLICANT: Ballinger, Dennis G.
APPLICANT: Pace, Ann M.
TITLE OF INVENTION: Interleukin-1 Hy2 Materials and Methods
FILE REFERENCE: 28110/35659
CURRENT APPLICATION NUMBER: US/09/316,081
CURRENT FILING DATE: 1999-05-20
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/08798414
Patent No. 6096728
GENERAL INFORMATION:
APPLICANT: COLLINS, David S.
APPLICANT: COLLINS, David S.
APPLICANT: GEVILACQUA, MICHAEL P.
TITLE OF INVENTION: COMPOSITION AND METHOD FOR TREATING
TITLE OF INVENTION: INFLAMMATORY DISEASES
    ; DB 4;
1.5e-31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  120 WFLCGPAEPQQPVQLTKESEPSA----RTKFYFEQ 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   120 WFLCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQ 154
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37.7%; Score 309; DB 45.2%; Pred. No. 1.5e-tive 23; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 1840 De Havilland Drive
CITY: Thousand Oaks
STATE: California
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/09316081
Patent No. 6339141
                                       Local Similarity 45.2% tes 70; Conservative
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CORRESPONDENCE ADDRESS:
ADDRESSEE: AMGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
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S-09-316-081-4
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| Batent No. 6294170 |
| GENERAL INFORMATION: |
| APPLICANT: Beciles and |
| APPLICANT: Collins and |
| APPLICANT: Coll
                                                                                                                                                                                                                                                                                                                68 CLSC---GTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCT 124
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                                                                                                                                                              10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSP--VILGVQGGSQ 67
                                                                                                                                                                                                                                       14 FRIWDVNOKTFYLRNNOLVAGYLQGPNVNLEEKIDVVP-----IEPHALFLGIHGGKM 66
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                                                                                 45; Indels 17;
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; Score 307.5; DB 4; Length 153;
; Pred. No. 2.3e-31;
16; Mismatches 45; Indels 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          125 SPEADQPVRLTQIPEDPAWDAPITDFYFQQ 154
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125 AMEADQPVSLINMPDE---GVMVTKFYFQE 151
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Fatent No. 5739282
GENERAL INFORMATION:
APPLICANT: COLOTTA, Francesco
APPLICANT: MAXTOVANT, Alberto
TILLE OF INVENTION:
ILL1 ANTAGONIST
NUMBER OF SEQUENCES: 113
CORRESPONDENCE ADDRESS:
    37.5%;
                                        Local Similarity 48.09
les 72; Conservative
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ORGANISM: Human
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US-08-476-860-10
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                                                Best Loca
Matches
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68 CLSC---GTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCT 124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 156;
                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
ADDRESSEE: BROWDY AND NEIMARK
STREET: 419 Seventh Street, N.W., Suite 300
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .4e-31
                                                                                                                                                                                                                                              CURRENT APPLICATION TALEGRAS #1.0, VEISION APPLICATION NUMBER: US/08/476,860 ETLING DATE: 07-7UN-1995 CLASSIFICATION: 435 PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: IT MI 94 A 002097 FILING DATE: 13-0CT-1994 ATTONNEY/AGGNT INFORMATION: NAME: BROWDY, ROGER L. REGISTRATION NUMBER: 25,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 37.5%; Score 307.5; Best Local Similarity 48.0%; Pred. No. 2.4¢ Matches 72; Conservative 16; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               125 SPEADQPVRLTQIPEDPAWDAPITDFYFQQ 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: COLOTTA-1
TELECOMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
                                                                                                           ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC COMPATIBLE
COMPUTER: DAMPER: PC DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Search completed: May 22, 2002, 14:09:22 Job time: 209 sec
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TELEX: 248633
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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May 22, 2002, 14:08:13 ; Search time 16.28 Seconds (without alignments) 920.758 Million cell updates/sec
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1 MMYLSGALCFRMKDSALKVL......IPEDPAWDAPITDFYFQQCD 156
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                       Title:
Perfect score:
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                                                                                                             Run on:
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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries pirl:* pir2:* pir3:* pir4:* PIR_71:* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	Score	3	319.5	307.5	307.5	297.5	289.5	137.5	134.5	129	128.5	128	127	122	119	84	84	84	80	80	78	78	76.5	76.5	75.5	75.5		7	73.5	73
	Result No.	п	7	m	4	S	9	7	ω,	o	10	11	12	13	14	12	16	17	18	19	20	21	22	23	24	25	56	27	28	29

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dnaK-type molecula	probable nelicase	probable serine/th	folate-dependent p	dihydropteroate sy	hypothetical prote	hypothetical prote	interleukin-1 alph	interleukin-1 alph	DnaJ protein (impo	dnaK-type molecula	probable proteinas	dnaK-tvpe molecula	dnaK-type molecula	dnaK-type molecula
T45471	1300/1 S26180	T01353	G97071	AF0902	A87374	D96554	ICB01A	A61246	AH2179	S37394	G70597	T05741	T03581	T04080
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632 2	1272 2	911	204	282	498	621	268	268	333	640	550	809	663	663
	8.9 1272 2							8.7 268 1	8.7 333	8.7 640	8.6 550	8.6 608	8.6 663	8.6 663
თ. თ დ. თ		8.8	8.7	8.7	8.7	8.7	8.7	8.7	71 8.7 333	8.7	70.5 8.6 550 ;	9.8	9.8	9.8

ALIGNMENTS

283138

Minimum DB seq length: 0 Maximum DB seq length: 200000000

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RESULT 1
J07104
J07104
J07104
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J07104
C; Date: 03-Dec-1999 #text_change 21-Jul-2000
C; Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 21-Jul-2000
B; Mullero, J.J.; Pace, A.M.; Nelken, S.T.; Loeb, D.B.; Correa, T.R.; Drmanac, R.; Ford Biochem. Biophys. Res. Commun. 263, 702-706, 1999
A; Title: ILLHYI: A novel interleukin-1 receptor antagonist gene.
A; Reference number: JC7104; MuID:99443727
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: 1-155 cMUL>
A; Cross references: GB:AF186094; NID:96049804; PIDN:AAF02757.1; PID:96049805
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILG 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 155;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                89.6%; Score 734; DB 2; Length 15
91.0%; Pred. No. 1.3e-66;
tive 4; Mismatches 10; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 91.03
Matches 141; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          A;Gene: illhy1
A;Map position: 2q14
C;Keywords: macrophage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          g
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RESULT A44610

interleukin-1 receptor antagonist precursor - mouse
N.Alternate names: IL-1Ra
N.Alternate names: IL-1Ra
N.Alternate names: IL-1Ra
N.Alternate names: IL-1Ra
S.Species: Mus musculus (house mouse)
C.Species: Musculus (house mouse)
C.Accession: A44610; B40956; A49031; I56106; I52970
C.Accession: A44610; B40956; A49031; I56106; I52970
B.Jood 78, 616-623, 1991
A.Itle: Cloning and expression of murine interleukin-1 receptor antagonist in macrop A.Reference number: A44610; MUID:91316273
A.Accession: A44610
A.Accession: A44610
A.Accession: A44610
A.Residues: I-178 <MArx
A.Residues: I-178 <MArx
A.Residues: I-178 <MArx
A.Residues: I-178 <MArx
A.Cross-references: GB:M64404; NID:9198296; PIDN:AAA39277.1; PID:9198297

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A; Molecule type: mRNA
A; Residues: 1-177 < CAR>
A; Residues: 1-177 < CAR>
A; Cross-references: GB:x53296; NID:g32578; PIDN:CAA37386.1; PID:g32579
A; Cross-references: GB:x53296; NID:g32578; PIDN:CAA37386.1; PID:g32579
A; Note: parts of this sequence, including the amino end of the mature protein, were c: R; Elsenberg, S.P.; Evans, R.J.; Arend, W.P.; Verderber, E.; Brewer, M.T.; Hannum, C.H!
Nature 343, 341-346, 1990
A; Title: Primarry structure and functional expression from complementary DNA of a human A; Reference number: S08160; MUID:90136921
A; Accession: S08160
                                                                                A.Cross-references: GB: M63099; NID:9186385; PIDN:AAB41943.1; PID:9186386
R.Lennard, A.; Gorman, P.; Carrier, M.; Griffiths, S.; Scotney, H.; Sheer, D.; Solari. Cytokine 4, 83-89, 1992
A.Tille: Cloning and chromsome mapping of the human interleukin-1 receptor antagonis: A; Reference number: 137894; MUID:92338323
A; Scatus: translated from GB/EMBL/DDBJ
A; Molecule type: DNA
A; Residues: 1-177 < LEN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: EMBL:X64532; NID:g33798; PIDN:CAA45832.1; PID:g33799
R;Carter, D.B.; Delbel Jr., M.R.; Dunn, C.J.; Tomich, C.S.C.; Laborde, A.L.; Slightom J.G.; Sleu, L.C.; Hardee, M.M.; Zurcher-Neely, H.A.; Reardon, I.M.; Heinrikson, R.L., Nature 344, 633-638, 1990
A;Title: Purification, cloning, expression and biological characterization of an inter A;Reference number: A30368; MUID:90220867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Status: not compared with conceptual translation
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Best Local Similarity 48.0%
Matches 72; Conservative
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                                                                                                                                                                             A/Accession: 1540956
BLILL: Desablid. T.E.; Taracey, D.E.; Blenkowski, M.J.
BLILL: J. Immunol. 21, 2775-2780, 1991
A/ATILL: J. Immunol. 21, 2775-2780, 1991
A/ATILL: J. Immunol. 21, 2775-2780, 1991
A/ACCESSION: A/4001

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C;Date: 07-Jun-1990 #sequence_revision 07-Jun-1990 #text_change 26-May-2000
C;Accession: A40956; 137894; A30368; S08160; S08159; A37822
R;Eisenberg, S.P.; Brewer, M.T.; Verderber, E.; Heimdal, P.; Brandhuber, B.J.; Thompson, Proc. Natl. Acad. Sci. U.S.A. 88, 5232-5236, 1991
A;Title: Interleukin 1 receptor antagonist is a member of the interleukin 1 gene family: A;Accession: A40956; MUID:91271363
                B.J.; Thompson
R;Elsenberg, S.P.; Brewer, M.T.; Verderber, E.; Heimdal, P.; Brandhuber, B.J.; Th
Proc. Natl. Acad. Sci. U.S.A. 88, 5232-5236, 1991
A;Title: Interleukin 1 receptor antagonist is a member of the interleukin 1 gene
A;Reference number: A40956; MUID:91271363
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGVQGGSQCL 69
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| DRPVSLINTPEEP---LIVTKFYFQE 176
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Best Local S:
Matches 71
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A; Retauts: not compared with conceptual translation
A; Residues: 1-177 < EL2>
A; Ribbard & Milcox, C.J.; Arend, W.P.; Joslin, F.G.; Dripps, D.J.; Helmdal, P.L.;
R; Hannum, C.H.; Wilcox, C.J.; Arend, W.P.; Joslin, F.G.; Dripps, D.J.; Helmdal, P.L.;
R; Hannum, C.H.; Milcox, C.J.; Arend, W.P.; Joslin, F.G.; Dripps, D.J.; Helmdal, P.L.;
A; Reference number: S08159; MUID:90136920
A; Molecule type: protein
A; Residues: 26-75;97-108; 110-116;120-131;163-176 < HAN>
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A; Residues: 26-57;97-108; 110-116;120-131;163-176 < HAN>
A; Residues: 26-52;97-108; 110-116;120-131;163-176 < HAN>
A; Residues: 26-52;97-108; 110-116;120-131;163-176 < HAN>
A; Residues: 26-52;97-108; 110-116;120-131;163-176 < HAN-Stimulated THP-1 cells
A; Residues: 26-52;97-108; 110-116;120-115; CAIE>
A; Molecule type: protein
A; Residues: 26-52;97-108; 110-116;120-115; CC Comment: For an alternative splice form, see PIR:A39386
C; Comment: For an alternative splice form, see PIR:A39386
C; Comment: For an alternative splice form, see PIR:A39386
C; Comment: Ror an alternative splice form, see PIR:A39386
C; Comment: Sold: 106/3
A; Residues: 20-52;70-77;122-127;170-175
A; Introns: 39/2; 69/1; 106/3
C; Superfamily: interleukin-1
C; Keywords: alternative splicing; cytokine receptor; extracellular protein; glycoprot F; 1-25/Domain: signal sequence #status predicted < SIG>
F; 26-177/Product: interleukin-1 receptor antagonist #status experimental
E; 109/Binding site: carbohydrate (Asn) (covalent) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'n
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSP--VILGVQGGSQ 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17;
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48.0%; Pred. No. 1.5e-23;
tive 16; Mismatches 45
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g

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Interleukin-1 receptor antagonist secreted form precursor - rabbit
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Species: Officer: 1964 # sequence_revision 18-Nov-1994 # text_change 16-Jul-1999
C;Accession: A44377; 146729
B;Cominelli, F; Bortolami, M.; Pizarro, T.T; Monsacchi, L; Ferretti, M.; Brewer, M.
J; Biol. Chem. 269, 6692-6971, 1994
A;Title: Rabbit interleukin-1 receptor antagonist. Cloning, expression, functional ch.
A;Residues: 1-177 < COM>
A;Nolecule type: mRNA
A;Residues: 1-177 < COM>
A;Note: sequence extracted from NCBI backbone (NCBIN:144169)
A;Experimental source: colon tissue
A;Note: sequence extracted from NCBI backbone (NCBIN:144169)
A;Rocession: AffOrd, K;;Myata, T; Ohkawara, S; Takao, T; Mori, S; Furukawa, S; Ma
Immunology 77, 235-244, 1992
A;Title: Interleukin-1 recetor antagonist in inflammatory exudate cells of rabbits. P
A;Accession: 146729
A;Status: translated from GB/EMBL/DDBJ
A;Accession: 146729
A;Status: translated from GB/EMBL/DDBJ
A;Accession: 1-177 < COT>
A;Cooss-references: GB:D21832; NID:9425787; PIDN:BAA04860.1; PID:9452205
C;Superfamily: interleukin-1
C;Keywords: cytckine receptor; extracellular protein; glycoprotein
F;1-25/Domain: signal sequence #status predicted <sigo
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C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Date: 08-Jun-1994 #sequence_revision 22-Nov-1996 #text_change 15-Oct-1999
C;Accession: S23010; S43047; S13092; B61246
                                                                                                                                                                                                                                                                                                                                                  70 SCGTEKG--PILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTSPE 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           68 CLSCGTEKGPILK--LEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTS 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                       39 FRIWDINOKTFYLRNNOLIAGYLOGPNTKLEEKIDMVP----IDFRNVFLGIHGGKLCL 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSP--VILGVQGGSQ 67
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                                                                                                                                                                         10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGVQGGSQCL
                                                Length 178;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 177;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 2;
                                                DB 2;
                                         36.3%; Score 297.5; DB 2
46.3%; Pred. No. 1.5e-22;
11ve 17; Mismatches 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            35.3%; Score 289.5; DB 2
46.3%; Pred. No. 9.7e-22;
tive 17; Mismatches 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     126 PEADQPVRLTQIPEDPAWDAPITDFYFQQ 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 153 ADHPVSLTNTPKEP---CTVTKFYFQE 176
                                                                                                                                                                                                                                                                                                                                                                                                                                        128 ADQPVRLTQIPEDPAWDAPITDFYFQQ 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           interleukin-1 beta precursor - sheep
                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 46.3
nes 69; Conservative
                                                                           Local Similarity
nes 68; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Si
Matches 69,
                                                Query Match
                                                                           Best Loc
Matches
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                                                                                                                                                                                                                                                                                    C. Accession: 137893; A39386

R. Muzio, M.; Polentarutti, N.; Sironi, M.; Poli, G.; De Gioia, L.; Introna, M.; Mantovan J. Sequence, Evision 11. Apr. 180.

R. Muzio, M.; Polentarutti, N.; Sironi, M.; Poli, G.; De Gioia, L.; Introna, M.; Mantovan J. Title: Cloning and characterization of a new isoform of the interleukin I receptor and A. Reference number: 137893; MUID:9535865

A. Reference number: 137893

A. Molecule type: mRNA
A. Residues: 1-180 c. RES>
A. Molecule type: mRNA
A. Residues: 1-180 c. RES>
A. Markin, G.; Van Le, L.; Morris, J.; Peace, A.; Bigler, C.F.; Jaffe, G.J.; P. C. Natl. Acad. Sci. U.S.A. 88, 3881-3885, 1991
A. Title: cDNA cloning of an intracellular form of the human interleukin I receptor antagan A. Reference number: A39386; MUID:9129436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Keywords: alternative splicing; cytokine receptor F;1-180/Product: interleukin-1 receptor antagonist, long intracellular splice form #stat F;1-3,25-180/Product: interleukin-1 receptor antagonist, short intracellular splice form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RiEisenberg, S.P.; Brewer, M.T.; Verderber, E.; Heimdal, P.; Brandhuber, B.J.; Thompson, Proc. Natl. Acad. Sci. U.S.A. 88, 5232-5236, 1991
A;Title: Interleukin I receptor antagonist is a member of the interleukin I gene family: A;Reference number: A40956; MUID:91271363
A;Accession: C40956
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ŝ
                                                                                                                                                           interleukin-1 receptor antagonist, long intracellular splice form - human N:Contains: interleukin-1 receptor antagonist, short intracellular splice form C; Species: Homo sapiens (man)
C; Species: 28-Feb-1992 *sequence_revision 11-Apr-1997 *text_change 26-May-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Interleukin-1 receptor antagonist precursor - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 20-Mar-1992 #sequence_revision 20-Mar-1992 #text_change 16-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLSC---GTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCT 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSP--VILGVQGGSQ 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Residues: 1-178 <EIS>
A; Rossidues: 1-178 <EIS>
A; Cross-references: GB:M63101; NID:g204928; PIDN:AAA41434.1; PID:g204929
C; Superfamily: interleukin-1
C; Keywords: cytokine receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A.Cross-references: GB:M55646; NID:9186291; PIDN:AAAS9138.1; PID:9186292
C.Comment: For an alternative splice form, see PIR:A30368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Score 307.5; DB 2;
; Pred. No. 1.5e-23;
16; Mismatches 45;
149 AMEADQPVSLTNMPDE---GVMVTKFYFQE 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          : |||||| || :|:: :| || |||:|: AMEADQPVSLTNMPDE---GVMVTKFYFQE 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    125 SPEADQPVRLTQIPEDPAWDAPITDFYFQQ 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: GDB:125897; OMIM:147679
A;Map position: 2q14.2-2q14.2
C;Superfamily: interleukin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     37.5%;
48.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: mRNA
A; Residues: 1-3,25-180 <HAS>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 72; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Gene: GDB: IL1RN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10
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A;Cross-references: GB:M37211; NID:g163200; PIDN:AAA30584.1; PID:g163201
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Matches 47; Conserv
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                                                                                        A;Accession: B61246
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-144,'L',146-266 <AND>
C;Comment: This protein lacks a conventional signal sequence for protein export. Cleavag ved form of interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
C;Comment: Interleukin-lbeta precursor is less heavily myristoylated than interleukin-la
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               interleukin-1 beta precursor - bovine

N;Alternate names: hematopoietin-1; IL-1 beta
C;Species: Bos primigenius taurus (cattle)
C;Date: 31-Mar-1989 #sequence_revision 31-Mar-1989 #text_change 22-Jun-1999
C;Accession: JL0010; S01380
R;Mallszewski, C.R.; Baker, P.E.; Schoenborn, M.A.; Davis, B.S.; Cosman, D.; Gillis, S.; A;Mallszewski, C.R.; Baker, B.E.; Schoenborn, M.A.; Davis, B.S.; Cosman, D.; Gillis, S.; A;Title: Cloning, sequence and expression of bovine interleukin 1-alpha and interleukin A;Reference number: A94695; MUID:88318652
A;Accession: JL0010
A;Accession: JL0010
A;Accession: JL0010
A;Residues: 1-266 cMAL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Superfamily: interleukin-1
C;Keywords: cytokine; immunoregulation; inflammation; lymphokine; macrophage; mitogen
F;114-266/Product: interleukin-1 beta #status predicted <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5
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R;Seow, H.F.; Rothel, J.S.; David, M.J.; Wood, P.R.
MAS Seq. 1, 423-426, 1991
A;Title: Nucleotide sequence of ovine macrophage interleukin-1 beta cDNA.
A;Reference number: S23010; MUID:92119335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17 LKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGVQGGSQCLSCGTEKG 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 266;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16.8%; Score 137.5; DB 1
30.9%; Pred. No. 3.3e-06;
11ve 24; Mismatches 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ouery Match
Best Local Similarity 30.93
Matches 42; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            135 TQIPEDPAWDAPITDF 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    248 GRF----RGGODITDF
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R: Leong, S.R.; Flaggs, G.M.; Lawman, M.; Gray, P.W.
Nucleic Acids Res. 16, 9054, 1988
A:Title: The nucleotide sequence for the cDNA of bovine interleukin-1 beta.
A:Reference number: S01380; MUID:89016591
A:Rolecule type: MRNA
A:Molecule type: MRNA
A:Rolecule type: MRNA
A:Residues: 1-251, A', 253-266 < LEDO>
A;Cross-references: RMBL:X12498; NID:9448; PIDN:CAA31018.1; PID:9449
C;Comment: This protein is a cytokine that mediates a variety of immunoregulatory and C;Comment: This protein lacks a conventional signal sequence for protein export. Clea
                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Comment: Interleukin-lbeta precursor is less heavily myristoylated than interleukin C;Superfamily: interleukin-1 C;Keywords: cytokine; immunoregulation; inflammation; lymphokine; macrophage; mitogen F;114-266/Product: interleukin-1 beta #status predicted <MAT>
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A;Accession: JN0724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A Molecule type: MRNA
A; Residues: 1-267 <HUE>
A; Residues: 1-267 <HUE>
A; Comment: 1-267 <HUE>
A; Comment: This protein is a pleiotropic cytokine that mediates a variety of processe c; Comment: This protein lacks a conventional signal sequence for protein export. Clear ved form of interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
C; Comment: Interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
C; Comment: Interleukin-lbeta precursor is less heavily myristoylated than interleukin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C.Keywords: cytokine; immunoregulation; inflammation; lipoprotein; lymphokine; macrop F;115-267/Product: interleukin-1 beta #status predicted <ILL>F;77/Binding site: myristate (Lys) (covalent) #status predicted
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C;Species: Sus scrofa domestica (domestic pig)
C;Date: 14-Jul-1994 #sequence_revision 22-Nov-1996 #text_change 22-Jun-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  77 --PILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTSPEADQPVRL 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           139 LKALHLLSQEMNREVVFCMSFVQGEERD-----NKIPVALGIKDKNLYLSC-VKKG 188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     57 PVILGVQGGSQCLSC-GTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESA 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKV----IKGEEISVVPNRALDASLS 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17 LKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGVQGGSQCLSCGTEKG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 266;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                M.P.; Molitor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match 16.4%; Score 134.5; DB 1
Best Local Similarity 30.9%; Pred. No. 6.7e-06;
Matches 42; Conservative 23; Mismatches 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Accession: JN0724
R;Huether, M.J.; Lin, G.; Smith, D.M.; Murtaugh,
Gene 129, 285-289, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             interleukin-1 beta precursor - pig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               135 TQIPEDPAWDAPITDF 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GHF----RGGQDITDF 259
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Rivandenbroeck, K.; Fiten, P.; Beuken, E.; Martens, E.; Janssen, A.; van Damme, J.; O Eur. J. Blochem. 217, 45-52, 1993
A;Title: Gene sequence, CDNA construction, expression in Escherichia coll and genetic A;Reference number: S38373; MUID:94039070
A;Accession: S38373
                                                                                         inflammation; lymphokine; macrophage; mitogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Title: Genomic sequence for human prointerleukin 1 beta: possible evolution from a A;Reference number: A25542; MUID:87040762
A;Accession: A25542 'As Accession: A25542 'As A;Molecule type: DNA; mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               interleukin-1 beta precursor [validated] - human
NiAlternate names: hematopoietin-1; IL-1 beta
C;Species: Homo saplens (man)
C;Date: 28-Feb-1986 #sequence_revision 15-May-1998 #text_change 15-Sep-2000
C;Accession: A25542; A29019; A404023; A93361; I51852; I65200; I38132; B27616; A01848;
R;Clark, B.D; Collins, K.L.; Gandy, M.S.; Webb, A.C.; Auron, P.E.
                                                                                                                                                                                                                                                                                                     9
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8
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C;Species: Sus scrofa domestica (domestic pig)
C;Date: 20-May-1994 #sequence_revision 01-bec-1995 #text_change 16-Jul-1999
C;Accession: S38373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Residues: 1-267 <VAN>
A;Cross-references: EMBL:X74568; NID:g407899; PIDN:CAA52660.1; PID:g407900
C;Genetics:
                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                        123 LHYRLRDEQQKSLVLSDPYELK-ALH----LNGQNI----NQQVIFSMSFVQGEPSNDKI 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PVILGVQGGSQCLSCGTEKG-PILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESA 115
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          62 VQGGSQCLSC-GTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 120
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                                                                                                                                                                                                                                                                                                         22;
                                                                              C; Keywords: cytokine; immunoregulation; inflammation; lymphokine F;118-269/Product: interleukin-1 beta #status experimental <ILL>
                                                                                                                                                                                                                               Length 269
                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                     LCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLS-
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                                                                                                                                                                                                                           15.6%; Score 128; DB 1; L 31.7%; Pred. No. 3.1e-05; live 23; Mismatches 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15.5%; Score 127; DB 2; 32.9%; Pred. No. 3.8e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        116 AYPGWFLCTSPEADQPVRL 134
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233 EFPNWYISTSQAEHKPVFL 251
                                                                                                                                                                                                                           Query Match 15.6%;
Best Local Similarity 31.7%;
Matches 44; Conservative
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                                                    C; Superfamily: interleukin-1
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Best Local Similarity
Matches 50; Conservat
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A; Residues: 1-267 <VAN>
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           IL-1-beta
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C:Superfamily: in
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Accession: S13029
Accession: S1
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A;Residues: 1-214 <KAT>
A;Residues: 1-214 <KAT>
A;Residues: 1-214 <KAT>
A;Cross-references: DDBJ:D42165; NID:92463549; PIDN:BAA22528.1; PID:92463550
A;Coomment: This protein mediates a variety of physiological response to infections and inthesis by hepatocytes, and stimulation of chondrocytes and synovial cells to produce synorial.y: interleukin-1
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                                                                                                                                                                                                                                      interleukin-1 beta - horse
C;Species: Equus caballus (domestic horse)
C;Date: 28-Oct-1997 #sequence_revision 28-Oct-1997 #text_change 20-Jun-2000
C;Accession: JC5646
R;Kato, H: Youn, H:Y.; Ohashi, T.; Watari, T.; Goitsuka, R.; Tsujimoto, H.; Hasegawa, Gene 177, 11-16, 1996
A;Title: Identification of an alternatively spliced transcript of equine interleukin-1 A;Reference number: JC5646; MUID:97080493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       interleukin-1 beta precursor - mouse
Nalternate names: hematopoletin-1; IL-1 beta
CSpecies: Mus mansculus (house mouse)
C;Dectes: Mus musculus (house mouse)
C;Date: 26-Jul-1996 #sequence_revision 22-Nov-1996 #text_change 22-Jun-1999
C;Date: 26-Jul-1996 #sequence_revision 22-Nov-1996 #text_change 22-Jun-1999
C;Accession: 155569; A24719; S13029
J. Immunol. 137, 3644-3648, 1986
A;Title: Two interleukin 1 genes in the mouse: Cloning and expression of the CDNA for Accession: 155969; MUID:87058957
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15.7%; Score 128.5; DB 2 33.6%; Pred. No. 2.1e-05; Live 16; Mismatches 44
116 AYPGWFLCTSPEADQPVRL--TQIPEDPAWDAPITDF 150
                                        230 LYPNWYISTSQAEQKPVFLGNSKGRQD-----ITDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    40; Conservative
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Best Local Similarity
Matches 40; Conserv
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C;Genetics:
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residues 117-269

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Acontents: annotation; Contouration, A.M.

Biochemistry 30, 2315-2323, 1991

Aritile: High-resolution three-dimensional structure of interleukin lbeta in solution A; Feference number: A44675; MUID: 91159409

A; Contents: annotation; (1)H-NMR structural determination
A; Feference number: A44675; MUID: 91159409

A; Contents: annotation; (1)H-NMR structural determination
A; Chem. 266, 7081-7086, 1991

A; Title: Structure-function mapping of interleukin 1 precursors. Cleavage leads to a A; Reference number: A39774; MUID: 91201363
A; Contents: annotation mapping of interleukin 1 precursors. Cleavage leads to a A; Reference number: A39774; MUID: 91201363
A; Contents: annotation
B; Finzel, B.C.; Matenpaugh, K.D.; Einspahr, H.M.
Submitted to the Brookhaven Protein Data Bank, December 1989
A; Reference number: A50016; PDB: IIIB
A; Contents: annotation; X-ray crystallography, 2.0 angstroms, residues 119-269
B; Finzel, B.C.; Clancy, L.L.; Holland, D.R.; Muchmore, S.W.; Watenpaugh, K.D.; Einspahr, J. Mol. Biol. 209, 779-791, 1980
A; Title: Crystal structure of recombinant human interleukin-lbeta at 2.0 angstrom reshifted to the anotation; X-ray crystallography, 2.0 angstroms
C; Comment: Ains protein lacks a conventional signal sequence for protein export. Cleaved form of interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
C; Comment: Interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
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N.Alternate names: hematopoietin-1; IL-1 beta; lymphocyte proliferation potentiating cispecies: Oryctolagus cuniculus (domestic rabbit)

C; Species: Oryctolagus cuniculus (domestic rabbit)

C; Date: 25-May-1989 #sequence_revision 22-Nov-1996 #text_change 22-Jun-1999

C; Accession: A27714; A30584; JU0082; A32166

E; Morchem. Biophys. Res. Commun. 150, 1237-1243, 1988

A; Riborl. S.; Goto, F; Goto, K; Ohkawara, S; Maeda, S.; Shimada, K.; Yoshinaga, M.

Biochem. Biophys. Res. Commun. 150, 1237-1243, 1988

A; Rifle: Cloning and sequence analysis of a cDNA for lymphocyte proliferation potentl A; Reference number: A27714; MUID:88134238

A; Accession: A27714

A; Accession: A27714

A; Molecule type: mRNA

A; Molecule type: mRNA

A; Residues: 1-268 «MOR>

B; Cannon, J.G.; Clark, B.D.; Wingfield, P.; Schmeissner, U.; Losberger, C.; Dinarello J. Immunol. 142, 2299-2306, 1989
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A; Map position: 2q13-2q21
A; Introns: 16/2; 33/3; 101/1; 156/1; 199/3
C; Superfamily: interleukin-1
C; Keywords: cytckine; immunoregulation; inflammation; lipoprotein; lymphokine;.macrop F; 17-69/Product: interleukin-1 beta #status experimental - TL1>
F; 76/Binding site: myristate (Lys) (covalent) #status experimental
F; 76/Binding site: carbohydrate (Asn) (covalent) #status absent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  62 VQGGSQCLSCG-TEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 120
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111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 111 | 1
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                                                    A; Note: modification of Arg-120 by phenylglyoxal blocks receptor binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 269;
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Best Local Similarity 29.0%; Pred. No. 0.00012;
Matches 45; Conservative 25; Mismatches 59; Indels
                                                                                                                       R;Clore, G.M.; Gronenborn, A.M.
submitted to the Brookhaven Protein Data Bank, January 1991
A;Reference number: A50049; PDB:Gilb
A;Contents: annotation; conformation by (13)C- and (1)H-NMR,
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A.; Residues: 1-5, KY, 7-269 cCLA>
A.; Cross-references: Gaing Cool And Tor 6-Lys, the DNA sequence had GaG for 6-Glu
R.; Mores are references: Gaing Cool And Tor 6-Lys, the DNA sequence had GaG for 6-Glu
R.; Mores are commander: A sequence had cool And Tor 6-Lys, the DNA sequence had GaG for 6-Glu
R.; Mores are commander: A sequence had cool And Tor 7: Residues: 1-26; Manuschi 
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A; Status: translated from GB/EMBL/DDBJ
A; Molecule type: mRNA
A; Residues: 1-269
A; Cross-references: EMBL:X56087; NID:g35662; PIDN:CAA39567.1; PID:g35663
A; Cross-references: EMBL:X56087; NID:g35662; PIDN:CAA39567.1; PID:g35663
B; Cabbo, K.M.; Wypych, J.; Yuschenkoff, V.N.; Lu, H.; Hunt, P.; Dukes, P.P.; Langley, K.B; Stebo, K.M.; Wypych, J.; Yuschenkoff, V.N.; Lu, H.; Hunt, P.; Dukes, P.P.; Langley, K.B; Stebo, K.M.; Mult:g8184226
A; Residues: 117-123, X', 125-126, X', 128 <2SE>
A; Residues: 118, 25-36, 1991
A; Title: The 31-kDa precursor of interleukin lalpha is myristoylated on specific lysines A; Reference number: A68293; Multip: 318056
A; Contents: annotation: myristylation of lysines
A; Title: The role of arginine residues in interleukin l receptor binding.
A; Title: The role of arginine residues in interleukin l receptor binding.
A; Title: The role of arginine residues in interleukin site
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151 AICIDNFS-----EGMSIGELTLDEERKNA---KRRTLGWTSLIGLSLFVSAVAGWF 199
                                                                                                        66 SQCLSCGTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTS----SFESAAYPGWF
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R; Finan, T.M.; Weidner, S.; Wong, K.; Buhrmester, J.; Chain, P.; Vorholter, F.J.; Hernan A; Fille: The complete sequence of the 1,683-kb psymB megaplasmid from the N2-fixing end A; Title: The complete sequence of the 1,683-kb psymB megaplasmid from the N2-fixing end A; Reference number: A95842; MUID:21396508; PMID:11481431

A; Reference number: A95842; MUID:21396508; PMID:11481431

A; Residues: 1-259 ckmRs

A; Residues: 1-250 ckmRs

A; Residues
                                                                               Title: Rabbit IL-1. Cloning, expression, biologic properties, and transcription during Reference number: A30584; MUID:89176242
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Best Local Similarity 27.2
Matches 41; Conservative
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14 DSALKVLYLH------NNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGVQGG 65

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us-09-770-528-2.rpr

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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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OM protein - protein search, using sw model

Run on:

May 22, 2002, 14:09:23 ; Search time 11.89 Seconds (without alignments) 508.011 Million cell updates/sec

Title: Perfect score: Sequence:

US-09-770-528-2 819 1 MMVLSGALCFRMKDSALKVL.....IPEDPAWDAPITDFYFQQCD 156

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

105224 tal number of hits satisfying chosen parameters:

105224 segs, 38719550 residues

Searched:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

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SwissProt_40:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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333 337 433 40 40	4444 4443 5443

ALIGNMENTS

RESULT 1 ILIX MOUSE	ID ILIX_MOUSE STANDARD; PRT; 178 AA.	J/; (Rel.	DT 01-WAY-1992 (Rel. 22, Last sequence update)	Interleukin-1 rece	(IRAP).	GN TELKN OR ID-IRA. OS Mus musculus (Mouse)	Eukaryota; Metazoa;	Mammalia, Eutheria, Rodentia,		RP SEQUENCE FROM N.A. (ISOFORM 1).	ra maduine=91230/12; Pudmed=1828262; RA Zahedi K., Seldin M.F., Rits M., Ezekowitz R.A., Whitehead A.S.,			7. C. TRUITUIO 1. 140.4220-4230 [1991].		MEDLINE=91316273; PubMed=1830498;	RA Matsushime H., Roussel M.F., Matsushima K., Hishinuma A., Sherr C.J.; RT "Cloning and expression of mirring interloubter reconter antendate		Blood 78:616-623(1991).			nc Sirainovals); RX MEDIJNE=447191; Duhmad=8001626.	Zahedi K.A., Uhlar C.M., Rits M.,	RT "The mouse interleukin 1 receptor antagonist protein: gene structure	and regulation in vitro.";	RL Cytokine 6:1-9(1994).				kA Gabay C., Porter B., Fantuzzi G., Arend W.P.,	and protein expression of intrac	RT distribution of secreted and intracellular IL-1 receptor antagonist in		לב טינות במשנים במשנים אין (באשל). SN [5]		MEDLINE=91271363; PubMed=1828896;	KA Elsenberg S.P., Brewer M.T., Verderber E., Helmdal P., RA Brandhuhar B.T. Thomnson B.C.	"Interleukin	tokine c	[6]	. W	
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antagonist protein precursor (IL-1RA) (ICIL-

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Homo sapiens
This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the FWBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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MEICWGPYSHLISLLILLFH -> MA (IN ISOFORM
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                Shuck M.E., Eessalu T.E., Tracey D.E., Bienkowski M.J.; "Cloning, heterologous expression and characterization of murine interleukin 1 receptor antagonist protein."; Eur. J. Immunol. 21:775-2780(1991).
                                                                                                                                                                                                              -!- SUBCELLULAR LOCATION: SECRETED (ISOFORM 1) OR INTRACELLULAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 178;
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48.6%; Pred. No. 1.7e-25;
tive 17; Mismatches 49
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BY SIMILARITY.
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PRINTS; PR00264; INTERLEUKIN1.
PRODOm; PD002536; Interleukin_1; 1.
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154 DRPVSLTNTPEEP---LIVTKFYFQE 176
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MEDIZINE-9920807; Pubbed-213180;
MEDIZINE-9920807; Pubbed-213180;
Carter D.B., Delbel M.R., Jr., Dunn C.J., Tomich C.S.C., Laborde A.L.,
Slightom J.L., Berger A.E., Bienkowski M.J., Sun F.F., McEwan R.N.,
Harris P. K.W., Yem A.W., Waszak G.A., Chosay J.G., Sidu L.C.,
Harris P. K.W., Yem A.W., Waszak G.A., Chosay J.G., Sidu L.C.,
Truesdell S.E., Shelly J.A., Reardon I.M., Hainrikson R.L.,
Truesdell S.E., Shelly J.A., Eessalu T.E., Taylor B.M., Tracey D.E.;
"Purification, cloning, expression and biological characterization of
an interleukin-1 receptor antagonist protein.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Primary structure and functional expression from complementary DNA of a human interleukin-1 receptor antagonist.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eisenberg S.P., Evans R.J., Arend W.P., Verderber E., Brewer M.T., Hannum C.H., Thompson R.C.;
                                                                                                                          Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A. (ISOFORM 1).
MEDLINE-92338323; PubMed-1365987;
Lennard A., Gorman P., Carrier M., Griffiths S., Scotney H.,
Sheer D., Solari R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Muzio M., Polentarutti N., Sironi M., Poli G., De Giola L.,
Introna M., Mantovani A., Colotta F.;
                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A. (ISOFORMS 1 AND 3).
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MEDLINE-90136921; Pubmed-2137201;
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MEDLINE-91271363; PubMed-1828896;
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MEDLINE-95355865; Pubmed-7629520;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=90136920; PubMed=2137200;
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Cytokine 4:83-89(1992).
1RA) (IRAP) (IL-1RÑ).
IL1RN OR IL1RA.
                                                                                        (Human)
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EMBL, U65590; AAB92269.1; -
EMBL, AF043143; AAC39672.1; -
EMBL, A30368; A37822.
PIR, A37822, A37822.
PIR, S08150; S08150.
PIR, S08159; S08159.
PIR, A40956, A40956.
PIR, A39386; A39386.
PDB, IITN: 30-APR-94.
PDB, IITN: 15-OCT-94.
PDB, IITN: 17-FEB-95.
PDB, IILR; 07-FEB-95.
PDB, IILR; 07
                                                                                                                                                                                            EMBL, M63099; AB41943.1; EMBL, X52015; CAA36262.1; EMBL, X53296; CAA37386.1; EMBL, X64532; CAA45832.1; EMBL, U65590; AAB92270.1; EMBL, U65590; AAB92270.1; EMBL, U65590; AAB92270.1; EMBL, U65590; AAB92270.1;
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EMBL; M63099; AAB41943.1; -.
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Best Local Similarity
Matches 72; Conserv
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O77482;
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15-DEC-1998
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                                                                                                                                                                                                                                                                                                                                SEQUENCE OF 35-177 FROM N.A. (ISOFORM 4).
MEDLINE-98183404; PubMed-9514884;
Weissbach L., Tran K., Colquhoun S.A., Champliaud M.F., Towle C.A.;
"Detection of an interleukin-1 intracellular receptor antagonist mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          X-RAY CRYSTALLOGRAPHY (2.1 ANGSTROMS).
MEDLINE=95172072; PubMed=7867645;
Schreuder H.A., Rondeau J.-M., Tardif C., Soffientini A., Sarubbi E., Akeson A., Bowlin T.L., Yanofsky S., Barrett R.W.;
Refined crystal structure of the interleukin-1 receptor antagonist. Presence of a disulfide link and a cis-proline.";
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-I- FUNCTION: IL-1RA INHIBITS THE ACTIVITY OF IL-1 BY BINDING TO ITS RECEPTOR. IL-1RA HAS NO IL-1 LIKE ACTIVITY.
-I- SUBCELLULAR LOCATION: SECRETED (ISOFORM 1) OR INTRACELLULAR (ISOFORMS 2, 3 AND 4).
-I- ALTERNATIVE PRODUCTS: 4 isoforms; 1 (shown here), 2/icIL-1ra, 3/icIL-1ra type II and 4; are produced by alternative splicing.
-I- TISSUE SPECIFICITY: THE INTRACELLULAR FORM OF IL-1RA IS PREDOMINANTLY EXPRESSED IN EPITHELIAL CELLS.
-I- SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stockman B.J., Scahill T.A., Roy M., Ulrich E.L., Strakalaitis N.A., Brunner D.P., Yem A.W., Deibel M.R. Jr.; "Secondary structure and topology of interleukin-1 receptor antagonist protein determined by heteronuclear three-dimensional NMR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                X-RAY CRYSTALLOGRAPHY (2.0 ANGSTROMS).
MEDLINE-94230368; PubMed-8175703;
Vigers G.Pa., Caffes P., Evans R.J., Thompson R.C., Eisenberg S.P.,
Brandhuber B.J.;
                                                                                   MEDLINE-9035444; PubMed-2143761; Bienkowski M.J., Eessalu T.E., Berger A.E., Truesdell S.E., Bienkowski M.J., Eessalu T.E., Berger A.E., Truesdell S.E., Shelly J.A., Laborde A.L., Zurcher Neely H.A., Reardon I.M., Heinrikson R.L., Chosay J.G., Tracey D.E.; "Purification and characterization of interleukin 1 receptor level antagonist proteins from THP-1 cells."; J. Biol. Chem. 265:14505-14511(1990).
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MEDILINE-9422051; PubMed-8045306;
Stockman B.J., Scahll T.A., Strakalaitis N.A., Brunner D.P.,
Yem A.W., Deibel M.R. Jr.;
"Solution structure of human interleukin-1 receptor antagonist
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "X-ray structure of interleukin-1 receptor antagonist at 2.0-A resolution.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biochem. Biophys. Res. Commun. 244:91-95(1998).
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MEDLINE-92297633; PubMed-1534997;
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Nature 343:336-340(1990).
                                                            SEQUENCE OF 26-52.
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use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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MEICRGLRSHLITLLLELEHS -> MAL (IN
ISOPORM 2).
MEICRGLRSHLITLLLELEHS -> MALADLYEEG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INTERLEUKIN-1 RECEPTOR ANTAGONIST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 177;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROSITE: PS00253; INTERLEDUKIN 1; 1.
Glycoprotein; Signal; Alternative splicing; 3D-structure.
SIGNAL 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     D1690776A7394057 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GEDNADSK (IN ISOFORM 3).
MISSING (IN ISOFORM 4).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               37.5%; Score 307.5; DB 1
48.0%; Pred. No. 2.8e-24;
ilve 16; Mismatches 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    174 AA
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(Rel. 37, Last sequence update)
                                                            tities requires a license agreement (;
send an email to license@isb-sib.ch)
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149 AMEADQPVSLTNMPDE---GVMVTKFYFQE 175
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR000975; Interleukin_1.
Pfam; PF00340; IL1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRINTS: PR00264; INTERLEUKINI.
ProDom; PD002536; Interleukin_1; 1.
SMART; SM00125; IL1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
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Sus scrofa (Pig).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss institute of Bioinformatics and the BMIL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (see http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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15-DEC-1998 (Rel. 37, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Interleukin-1 receptor antagonist protein precursor (IL-1RA) (IL-1RN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLSCGTEKGP1-LKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTSP 126
                     Interleukin-1 receptor antagonist protein precursor (IL-1RA) (IL-1RN)
                                                                                                                                                                                                                              SEQUENCE FROM N.A.
MEDLINE-98305607; PubMed-9643454;
Kirisawa R., Fukuda T., Yamanaka H., Hagiwara K., Goto M., Obata Y.,
Yoshino T., Iwai H.;
"Enzymatic amplification and expression of bovine interleukin-1
                                                                                                                                                                                                                                                                                                                                                                 Vet. Immunol. Immunopathol. 62:197-208(1998).
-!- FUNCTION: IL-1RA INHIBITS THE ACTIVITY OF IL-1 BY BINDING TO ITS
RECEPTOR. IL-1RA HAS NO IL-1 LIKE ACTIVITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSP--VILGVQGGSQ 67
                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BY SIMILARITY.
N-LINKED (GLCNAC. .) (POTENTIAL).
1E56E7F224FF851F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INTERLEUKIN-1 RECEPTOR ANTAGONIST PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 174;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            36.8%; Score 301.5; DB 1
48.0%; Pred. No. 1.1e-23;
tive 15; Mismatches 49
15-JUL-1999 (Rel. 38, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      177 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                       -!- SIMILARITY: BELONGS TO THE IL-1 FAMILY.
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EADQPVGLINMPTEA---LKVTKFYFQQ 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EADQPVRLTQIPEDPAWDAPITDFYFQQ 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROSITE: PS00253; INTERLEUKIN_1; 1. Glycoprotein; Signal. 1. 23 BY SIM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; AB005148; BAA31854.1; -.
HSSP; P18510; 11LR.
InterPro; IPR000975; Interleukin_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pfam; PF00340; ILI; 1.
PRINTS; PR00240; ILI; INTERLEUKINI.
PRODOM; PD002536; INTERleukin_1; 1.
SMART; SM00125; III; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
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                                                                                                                                                                                                                                                                                                                                                   receptor antagonist cDNA.";
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107 1
174 AA;
                                                                                          Bos taurus (Bovine).
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hes 71; Conserv
                                                                                                                                                                                   NCBI_TaxID-9913;
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IL1RN OR IRAP1
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ID ILIX_PIG

AC 029056;

DT 15-DEC-199

DT 15-DEC-199

DT 15-UL-199

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GN ILIRN OR 1
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                                                  IRAP).
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Matches
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    SOUR PRESENTATION OF THE PRESENT OF 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Interleukin-1 receptor antagonist protein precursor (IL-1RA) (IL-1RN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                              38 FRIWDVNQKTFYLRNNQLVAGYLQGPNTKLEEKIDVVP-----VEPHFVFLGIHGGKL 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSP--VILGVQGGSQ 67
Euteleostomi;
Sus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BY SIMILARITY.
INTERLEUKIN-1 RECEPTOR ANTAGONIST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 177;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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2114DC6119A9D5F9 CRC64;
Chordata; Craniata; Vertebrata;
Cetartiodactyla; Suina; Suidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      36.6%; Score 299.5; DB 1
48.0%; Pred. No. 1.8e-23;
iive 15; Mismatches 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-NAY-1992 (Rel. 22, Created)
01-MAY-1992 (Rel. 22, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 178 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; L38849; AAA99424.1; -.
HSSP; P18510; 11RA.
InterPro; IPR000975; Interleukin_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRINTS; PR00264; INTERLEUKINI.
PRODOM; PR002536; Interleukin_1; 1.
SMART; SM00125; ILL; 1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
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SEQUENCE FROM N.A.
MEDLINE-91271353; PubMad-1828896;
                                                                                       SEQUENCE FROM N.A.
STRAIN-CROSSBREED; TISSUE-Lung;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20093 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 46.v...
Rest Local Similarity 72; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Glycoprotein; Signal
    Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           91
109
177 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pfam; PF00340; IL1;
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                                           NCBI_TaxID=9823
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CARBOHYD
SEQUENCE
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P25086;
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018999; 077745;
15-DEC-1998 (Rel. 37, Created)
15-DEC-1999 (Rel. 37, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Interleukin-1 réceptor antagonist protein precursor (IL-1RA) (IL-1RN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               68 CLSCGTEKGPILK -- LEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTS 125
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                                                                                                                                                                                                                                                                                   Goto F., Goto K., Miyata T., Ohkawara S., Takao T., Mori S., Purukawa S., Maeda T., Iwanaga S., Shimonishi Y., Yoshinaga M.; "Interleukin-I receptor antagonist in inflammatory exudate cells rabbits. Production, purification and determination of primary
                                                                                                                                                                                                                                                                                                                                                                                                                                                          -!- FUNCTION: IL-1RA INHIBITS THE ACTIVITY OF IL-1 BY BINDING TO RECEPTOR. IL-1RA HAS NO IL-1 LIKE ACTIVITY.
-!- SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BY SIMILARITY.
INTERLEUKIN-1 RECEPTOR ANTAGONIST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 177;
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                                                                                                                                    Hamada H., Mulligan R.C.;.
Submitted (XXX-1992) to the EMBL/GenBank/DDBJ databases.
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N-LINKED (GLCNAC. . .) (P
F5BC087F097FFEAF CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          35.3%; Score 289.5; DB 1; 46.3%; Pred. No. 1.9e-22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 126 PEADQPVRLTQIPEDPAWDAPITDFYFQQ 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         150 LEADQPVSLTNTPDD---SIVVTKFYFQE 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRINTS; PRO0264; INTERLEUKIN1.
PRODOM; PRO02536; Interleukin_1; 1.
SMARF; SM00125; ILL; 1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
               inflammation.";
J. Biol. Chem. 269:6962-6971(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HSSP; P18510; IILR.
InterPro; IPR000975; Interleukin_1.
Pfam; PF00340; ILL; 1.
                                                                                                                                                                                                                                                       MEDLINE-93052512; Pubmed-1427977;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20214 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; S68977; AAB30093.1; --
EMBL; M57526; AAA31374.1; --
EMBL; D21832; BAA04860.1; --
PIR; A54377; A54377.
                                                                                                                                                                                                                                                                                                                                                                                                                              Immunology 77:235-244(1992)
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109 1
177 AA;
                                                                                                         N.A.
                                                                                                                                                                                                  [3]
SEQUENCE FROM N.A.
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Matches 69; Conserv
                                                                                                         SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                        structure."
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CARBOHYD
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                     RRITH REPORT OF THE PRICE OF TH
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                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eisenberg S.P., Brewer M.T., Verderber E., Heimdal P.,
Brandhuber B.J., Thompson R.C.;
"Interleukin I receptor antagonist is a member of the interleukin I
gene family: evolution of a cytckine control mechanism.";
Proc. Natl. Acad. Sci. U.S.A. 88:5232-5236(1991).
-!- FUNCTION: IL-IRA INHIBITS THE ACTIVITY OF IL-1 BY BINDING TO ITS
RECEPTOR. IL-1RA HAS NO IL-1 LIKE ACTIVITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SCGTEKG--PILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTSPE 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-AUG-1992 (Rel. 23, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Interleukin-1 receptor antagonist protein precursor (IL-1RA) (IL-1RN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGVQGGSQCL 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Cominelli F., Bortolami M., Pizarro T.T., Monsacchi L., Ferretti M.,
Brewer M.T., Eisenberg S.P., Ng R.K.;
"Rabbit interleukin-1 receptor antagonist. Cloning, expression,
functional characterization, and regulation during intestinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oryctolagus cuniculus (Rabbit).
Eukaryota Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
NCBI_TaxID=9986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROTEIN.

S SIMILARITY.

N-IINKED (GLCHAC. . .) (POTENTIAL)
F3A5754FB6C51B03 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BY SIMILARITY.
INTERLEUKIN-1 RECEPTOR ANTAGONIST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      36.3%; Score 297.5; DB 1; Length 36.3%; Pred. No. 2.9e-23; 1ve 17; Mismatches 51; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             177 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; M63101; AAA41434.1; -.
PIR; C40956; C40956.
BSSP; P18510; 11LK
INTERPRO; IPR000975; INTERleukin_1.
PFam; PF00340; ILI.
PRINTS; PR00264; INTERLEUKINI.
PRODOM; PD002536; INTERLEUKIN.1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                153 ADHPVSLTNTPKEP---CTVTKFYFQE 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 128 ADQPVRLTQIPEDPAWDAPITDFYFQQ 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ERUSITE; PS00253; INTERLEUKIN_1; 1. Glycoprotein; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
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01-AUG-1992 (Rel. 23, Last sequ
15-DEC-1998 (Rel. 37, Last anno
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nes 68; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           142
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110 1
178 AA;
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ID ILIX_RABIT
AC P26890;
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CONFLICT
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-97366446; PubMed-9223227;
Kato H., Ohashi T., Matsushiro H., Watari T., Goitsuka R.,
Tsujimoto H., Hasegawa A.;
"Molecular cloning and functional expression of equine interleukin-1
                                                                                                                                                                                                                                                                                                                                                                   BY BINDING TO ITS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutherla; Perissodactyla; Equidae; Equus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INTERLEGKIN-1 RECEPTOR ANTAGONIST PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             32.9%; Score 269.5; DB 1; Length 177;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  51; Indels
                                                                                                                                                                                                                                                                         N-LINKED (GLCNAC. . .) (PORT | LABC377F1F1CF80B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Pred. No. 2.1e-20; 20; Mismatches 51
                                                                                                                                                                                                                     Vet. Immunol. Immunopathol. 56:221-231(1997).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BY SIMILARITY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR000975; Interleukin_1.
Pfam; PF00340; ILL; 1.
PRINTS; PR00264; INTERLEUKINI.
Procom; PD002536; Interleukin_1; 1.
SMART; SM00125; ILL; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PS00253; INTERLEUKIN_1; 1. cein; Signal.
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ID ILIB_SHEEP STANDARD;
AC P21621;
DT O1-MAY-1991 (Rel. 18, Created)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            U92482; AAC39257.1; -.
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                   Equus caballus (Horse).
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109
                                                                                                                                                                                                    receptor antagonist.";
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177 AA;
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                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                       NCBI_TaxID=9796;
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109
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OGNOOON SANARAN SANARA
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MEDLINE-91088326; PubMed-2263490;

REDLINE-91088326; PubMed-2263490;

RESCASTAND C. Sargan D.,

RESCASTAND C.,

RIVELOCIIDE Sequence of ovine interleukin-1 beta.";

Nucleic Acids Res. 18:7165-7165(1990).

LI-THAMOSYTE PROLIFERATION & FIRNOBLAST GROWTH FACTOR ACTIVITY.

THYMOSYTE PROLIFERATION, & FIRNOBLAST GROWTH FACTOR ACTIVITY.

THYMOSYTE PROLIFERATION BY INDUCING IL-2 RELEASE, BEING

IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING

IDENTIFIED AS ENDOGENOUS PROCEEDS.

THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE FROM SYNOVIAL CELLS.

THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.

AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.

THE SECRETAL BOUNCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS.

PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS.

SECRETARY PROFILEINS.

SECRETARY PROFILEINS.

THAT USED FOR OTHER
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                                                                                                                                                                                                                                                                                                                                  MEDLINE-92119335; PubMed-1840515; Seow H.F., Rothel J.S., David M.J., Wood P.R.; "Nucleotide sequence of ovine macrophage interleukin-1 beta cDNA."; DNA Seq. 1:423-426(1991).
                                                                                                                        Ovis aries (Sheep).
Sukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Caprinae; Ovis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17;
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PROSITE; PS00253; INTERLEUKIN_1; 1.
Cytokine; Macrophage; Mitogen; Inflammatory response; Pyrogen.
PROPEP 1 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 266;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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Q -> K (IN REF. 2).

V -> A (IN REF. 2).

P -> L (IN REF. 2).

P -> L (IN REF. 2).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 137.5; DB
Pred. No. 9.5e-07
01-MAR-1992 (Rel. 21, Last sequence update) 30-MAY-2000 (Rel. 39, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -! - SIMILARITY: BELONGS TO THE IL-1 FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; X54796; CAA38566.1; -.
EMBL; X56972; CAA40293.1; -.
PIR; S13092; S13092.
PIR; S131810; S13810.
PIR; S23010; S23010.
HSSP; P01584; 511B.
InterPro; IPR002348; ILL_HBGF.
InterPro; IPR000975; Interleukin_1_prop.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INTERLEUKIN-1
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                                                             Interleukin-1 beta precursor (IL-1 beta).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or send an email to license@isb-sib.ch).
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Pfam; PF02394; IL1_propep; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16.8%;
30.9%;
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30717 1
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Best Local Similarity
Matches 42; Conserv
                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                       NCBI_TaxID=9940;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SUBUNIT: MONOMER (BY SIMILARITY).
SUBUNIT: MONOMER (BY SIMILARITY).
SUBUNIT: MONOMER (BY SIMILARITY).
ALIERNATIVE PRODUCTS: 2 isoforms; a long form (shown here) and a short form; are produced by alternative splicing.
DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENUS OF THESEES PROPERINE SREWE SOME AS YET UNDEFINED FUNCTION.
MISCELLANBOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS.
OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kato H., Ohashi T., Nakamura N., Nishimura Y., Watari T., Goitsuka R.,
Tsujimoto H., Hasegawa A.;
"Molecular cloning of equine interleukin-1 alpha and ·beta cDNAs.";
Vet. Immunol. Immunopathol. 48:221-231(1995).
                                139 LKALHLPSQEMSREVVFCMSFVQGEERD------NKIPVALGIRDKNLYLSC-VKKG 188
                                                                     --PILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTSPEADQPVRL 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGVQGGSQCLSCGTEKG 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-98285941; PubMed-5622738; Howard R.D., McIlwraith C.W., Trotter G.W., Nyborg J.K.; Cloning of equine interleukin-1 alpha and equine interleukin-1 alpha and determination of their full-length cDNA sequences."; Am. J. Vet. Res. 59:704-711(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-97080493; Pubmed-8921838;
Kato H., Youn H.Y., Ohashi T., Watari T., Goitsuka R., Tsujimoto H.,
                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Identification of an alternatively spliced transcript of equine
                                                                                                                                                                                                                                                                                      028386; 077744; 018995;
15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
16-DEC-1998 (Rel. 31, Last sequence update)
Interleukin-1 beta precursor (IL-1 beta)
                                                                                                                                                                                                                                                                              268 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. (SHORT ISOFORM).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A. (LONG ISOFORM)
MEDLINE-96131982; Pubmed-8578682;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. (LONG ISOFORM)
                                                                                                                                              135 TQIPEDPAWDAPITDF 150
                                                                                                                                                                                  248 GRF----RGGODITDF 259
                                                                                                                                                                                                                                                                             STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                        Equus caballus (Horse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9796;
                                                                                                                                                                                                                                                                             L1B_HORSE
                                                                                                                                                                                                                                      RESULT 9
IL1B_HORSE
17
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(BY, SIMILARITY).
SUBUNIT: MONOMER (By similarity).
DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE
                                                                                                                                                                                                                                                                                                                                                                                                                                      133 LVLSGA------ETDKIPVALG 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 62 VQGGSQCLSCGTEKG-PILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                    61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MATURATION AND PROLIFERATION, AND FIBROBLAST GROWTH FACTOR ACTIVITY. IL-128 ARE INVOLVED IN THE INFAMMATORY RESPONSE, BEING IDENTIFIED AS ENDOGENOUS PROGENS, AND ARE REPORTED TO STIMILATE THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE FROM SYNOVIAL CELLS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Cervoidea; Cervidae; Cervinae; Cervus.
                                                                                                                                                                                                                                                                                                                                                                                                                  2 MYLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUBMITTED (MAR-1995) to the EMBL/GenBank/DDBJ databases.
-1- FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL
                                                                                                                                                                     PROSÍTE; PSO0253; INTERLEUKIN_1; 1.
Cytokine; Macrophage; Mitogen; Inflammatory response; Pyrogen;
                                                                                                                                                                                                                                                                                                                                                                      Length 268;
                                                                                                                                                                                                      BY SIMILARITY.
INTERLEUKIN-1 BETA.
MISSING (IN SHORT ISOFORM).
D -> N (IN REF. 2).
H -> Q (IN REF. 2).
AM -> VV (IN REF. 2).
V -> M (IN REF. 2).
TO -> N (IN REF. 2).
                                                                                                                                                                                                                                                                                          EG -> DD (IN REF. 2).
M -> V (IN REF. 2).
S -> K (IN REF. 2).
336F27792A1542EA CRC64;
                                                                                                                                                                                                                                                                                                                                                                  16.7%; Score 137; DB 1;
30.3%; Pred. No. 1.1e-06;
Live 22; Mismatches 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-0cT-1996 (Rel. 34, Created)
01-0cT-1996 (Rel. 34, Last sequence update)
01-MAR-2002 (Rel. 41, Last annotation update)
121 FLCTSPEADQPVRL--TQIPEDPAWDAPITDFYFQ 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        236 YISTSQAEKSPVFLGNTRGGRD-----ITDFIME 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      266 AA
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                                                                                                                                                                                                                                                                                                                                  MW.
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                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STANDARD;
                                                                                                                                                                                                                                                                                                                               268 AA;
                                                                                                                                                                                                                                                                                                                                                                               Similarity
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Best Local Simi
Matches 47;
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CONFLICT
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ILIB_CAVPO
ID ILIB_CA
AC Q9WVG1,
DT 01-MAR:
DT 01-MAR:
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                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its way non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5
AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.

-!- MISCELLANDOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGNENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER SECRETORY PROTEINS (By SIMILARITY).

-!- SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LEONG S.R., Flaggs G.M., Lawman M., Gray P.W.;
MEDLINE-89016591; Pubmed-3262866;
MEDLINE-89016591; Pubmed-3262866;
MEONG S.R., Flaggs G.M., Lawman M., Gray P.W.;
"The nucleotide sequence for the CDNA of bovine interleukin-1 beta.";
Nucleic Acids Res. 16:9054-9054(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               77 --PILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTSPEADQPVRL 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-88318652; Pubmed-3261832;
Maliszewski C.R., Baker P.E., Schoenborn M.A., Davis B.S., Cosman D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17 LKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGVQGGSQCLSCGTEKG 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17;
                                                                                                                                                                                                                                                                                                                                                      Cytokine; Macrophage; Mitogen; Inflammatory response; Pyrogen. PROPEP 1 113 BY SIMILARITY.
CHAIN 114 266 INTERLEUKIN-1 BETA.
SEQUENCE 266 AA; 30629 MW; ~4F40B4E6F0D9F060 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                       16.7%; Score 136.5; DB 1; Length 266; 30.9%; Pred. No. 1.2e-06; Live 23; Mismatches 54; Indels 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Interleukin-1 beta precursor (IL-1 beta)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 266 AA.
                                                                                                                                                                                                                                Interpro; IPR002348; ILL_HBGF.
Interpro; IPR000975; Interleukin_l.
Interpro; IPR003502; Interleukin_l_prop.
                                                                                                                                                                                                                                                                     Pfam; PF00340; III; 1.
Pfam; PF02394; III_propep; 1.
PRINTS; PR00262; IIIHBGF.
ProDom; PD002536; Interleukin_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                                                                                                                                                                                                              SMART; SM00125; IL1; 1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
                                                                                                                                                                                                       EMBL; U20500; AAA62234.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   135 TQIPEDPAWDAPITDF 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best_Local Similarity 30.9
Matches 42; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               248 GHF----RGGODITDF 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STANDARD;
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                                                                                                                                                                                                                                                                  -i- SUBUNIT: MONOMER.
-i- DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.
-i- MISCELLANEOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGNENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               interleukin 1 beta complementary DNAS.";
Mol. Immunol. 25:429-437(1988).
-!- FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES
- THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL
MATURATION & PROLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY.
IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING
IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE FROM SYNOVIAL CELLS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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'Cloning, sequence and expression of bovine interleukin 1 alpha and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 266;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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9D1EF8F575070586 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 54;
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(Rel. 41, Last sequence update)
(Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                              SECRETORY PROTEINS. SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23; Mismatches
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InterPro; IPR000975; Interleukin_1.
InterPro; IPR000975; Interleukin_1.
Pfam; PF00340; ILL; 1.
Pfam; PF00349; ILL_propep; 1.
PRINTS; PR00262; ILLHBGF.
SMART; SM00125; ILL; ILL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00253; INTERLEUKIN_1;
Cytokine; Macrophage; Mitogen; In
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; M35589; AAA30585.1; -.
EMBL; X12498; CAA31018.1; -.
EMBL; M37211; AAA30584.1; -.
PIR; JL0010; ICB01B.
PIR; S01380; S01380.
HSSP; P01584; IHIB.
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30774 MW;
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266 AA;
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hes 42; Conserv
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01-MAR-2002 (
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01-AUG-1992
01-AUG-1992
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Э
                                                                                                                                                                                                                                                                                                       (BY SIMILARITY).
SUBDNIT: MONOMER (BY SIMILARILY)
DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE
DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE
AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.
MISCELLANEOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE
PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS
OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
SECRETORY PROPEINS (BY SIMILARILY).
SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                expression
                                                                                                                                                                      Molecular cloning of the guinea pig GRO gene and its rapid expression in the tissues of lipopolysaccharide-injected guinea pigs.";
Int. Arch. Alleryy Immunol. 119:101-111(1999).
-1- FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL MATURATION AND FROLIFERATION, AND FIBROBLAST GROWTH FACTOR ACTIVITY. IL-1S ARE INVOLED IN THE INFLAMMATORY RESPONSE, BEING IDENTIFIED AS ENDOGENGY PYROGENS, AND ARE REPORTED TO STIMULATE THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE FROM SYNOVIAL CELLS COMMENTATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                140 LKALHLNGDNLNRQVVFSMSFVQGER----SDNKMPVALGLKGKNLYLSCVMKDG 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILGVQGGSQCLSCGTEKG 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             77 -PILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTSPEADQPVRL 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cavia porcellus (Guinea pig).
Wakaryota: Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Hystricognathi; Caviidae; Cavia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR000975; Interleukin_1.
InterPro; IPR003502; Interleukin_1_prop.
Pfam; PF00340; IL1, 1.
Pfam; PF00344; IL1_propep; 1.
PRINTS; PR00262; IL1HBGF.
SMART; SM00125; IL1, 1.
PROSITE; PS00253; INTERLUKIN_1; 1.
CYtokine; Macrophage; Mitogen; Inflammatory response; Pyrogen.
PROSITE; PS00253; INTERLUKIN_1; 1.
CYtokine; Macrophage; Mitogen; Inflammatory response; Pyrogen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 266;
                                                                                                                                              Yoshimura T., Takeya M., Ogata H., Yamashiro S., Modi W.S.,
Gillitzer R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INTERLEUKIN-1 BETA.
46558BA16B4C529A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
16.1%; Score 131.5; DB 1
Best Local Similarity 31.9%; Pred. No. 3.9e-06;
Matches 38; Conservative 21; Mismatches 45
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Interleukin-1 beta precursor (IL-1 beta).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        267
                                                                                                               STRAIN-2; TISSUE-Spleen;
MEDLINE-99323828; PubMed-10394101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; AF119622; AAD38502.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30530 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR002348; IL1_HBGF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P10749; 811B.
                                                                                      [1]
SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ILIB_PIG
ID ILIB_PIG
AC P26889;
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8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           57 PVILGVQGGSQCLSC-GTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESA 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R HSSP; P01584; IHIB.
R InterPro; IPR002348; ILL_HBGF.
InterPro; IPR003502; Interleukin_1.
R InterPro; IPR003502; Interleukin_1.
R Fām; PF00340; ILL]; Interleukin_1.
R Fām; PF00340; ILL]; Interleukin_1; I.
R RAMT; PR002536; Interleukin_1; I.
R RAMT; SR001255; ILL]; I.
R RAMT; SR001255; ILL]; I.
R RAMT; ROOU125; ILL]; I.
R Cytokine; Macrophage; Mitogen; Inflammatory response; Pyrogen.
R PROPEP
                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom1;
Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAEKV----IKGEEISVVPNRALDASLS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INTERLEUKIN-1 BETA.
7F6B92B784D5086F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        116 AYPGWFLCTSPEADQPVRL--TQIPEDPAWDAPITDF 150
01-AUG-1992 (Rel. 23, Created)
01-AUG-1992 (Rel. 23, Last sequence update)
03-MAY-2000 (Rel. 39, Last annotation update)
Interleukin-1 beta precursor (IL-1 beta).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15.8%; Score 129; DB 1.
29.9%; Pred. No. 7e-06;
tive 26; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; M86725; AAA02584.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      47; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SECRETORY PROTEINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PIR; JN0724; JN0724.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    267 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                             Sus scrofa (Pig)
                                                                                                                                                                              NCBI_TaxID=9823;
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Page 10

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InterPro; IPR003502; Interleukin_1_prop
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION. MISCELLANBOUGS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUBUNIT: MONOMER.
DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE
                                                                                                                                                                                                                                                                                                    Telford J.L., Macchia G., Massone A., Carinci V., Palla E., Melli M.; "The murine interleukin 1 beta gene: structure and evolution."; Nucleic Acids Res. 14:9955-9963(1986).
                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDINE-87058957; PubMed-3491144;
Gray P.W., Glaister D., Chen E., Goeddel D.V., Pennica D.;
"Two interleukin 1 genes in the mouse: cloning and expression of the CDNA for murine interleukin 1 beta.";
J. Immunol. 137:3644-3648(1986).
                                                                                                                                                                                                                                                                                                                                                     SEQUENCE OF 118-139.
MEDLINE-88229074; PubMed-2967326;
MEDLINE-80.2, Newton R.C., Rutledge S.J., Horuk R., Matthew J.B.,
Covington M., Lin Y.;
"Characterization of murine IL-1 beta. Isolation, expression, and
                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                            01-JUL-1989 (Rel. 11, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Interleukin-1 beta precursor (IL-1 beta).
                                       269 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Interpro; IPR002348; IL1_HBGF.
Interpro; IPR000975; Interleukin_1.
                                       PRT;
                                                                                                                                                                                                                                                                                          MEDLINE-87117546; PubMed-3492706;
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                                                                 (Rel. 11, Created)
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                                        STANDARD;
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PIR; S13029; S13029.
PDB; B11B; 15-OCT-94.
PDB; ZMIB; 31-JAN-94.
                                                                                                                              musculus (Mouse).
                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                  NCBI_TaxID=10090;
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                                                                01-JUL-1989
                                                                             01-JUL-1989
                                        IL1B_MOUSE
                           (L1B_MOUSE
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PVILGVQGGSQCLSCGTEKG-PILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESA 115
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Mammalia; Metatheria; Diprotodontia; Phalangeridae; Trichosurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE-99221044; PubMed-10206203; Wedlock D.N., Goh L.P., Parlane N.A., Buddle B.M.; "Molecular cloning and physiological effects of brushtail possum interleukin-lbgta."; .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22;
                                                            SMART; SM00125; IL1; 1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
Cytokine; Macrophage; Mitogen; Inflammatory response; Pyrogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15.6%; Score 128; DB 1; Length 269; 31.7%; Pred. No. 9e-06; tive 23; Mismatches 50; Indels
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                                                                                                                                 INTERLEUKIN-1 BETA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAR-2002 (Rel. 41, Created)
01-MAR-2002 (Rel. 41, Last sequence update)
01-MAR-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          269 AA.
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Pfam; PF00340; ILL; 1.
Pfam; PF02394; ILL_propep; 1.
PRINTS; PR00262; ILLHBGF.
PRODOM: PD002536; Interleukin_1; 1.
SWART; SM00125; ILL; 1.
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269 AA;
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Best Local Similarity
Matches 44; Conserv
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Vet. Immunol. Immunopathol. 67:359-372(1999).

-!- FUNCTION: PRODICED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES
THYMOCYTE PROLIFERATION BY INDICING IL-2 RELEASE, B-CELL
MATURATION & PROLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY.
IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING
IDENTIFIED AS ENROGENOUS PROGENS, AND ARE REPORTED TO STIMULATE
THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE FROM SYNOVIAL CELLS.
-!- DOMAINT: MONOMER.
-!- DOMAINT THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE
AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.
-!- MISCELLANEOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGRENT IN THE
PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS
ONE IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
-!- SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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InterPro; ...

InterPro; ...

Ream; PF00340; ILL; 1.

Ream; PF00340; ILL; 1.

Ream; PF002540; ILL]

PF000125; ILL; 1.

Ream; PR00125; ILL; 1.

Ream; PR00125; ILL; 1.

Ream; PF00125; I
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InterPro; IPR002348; IL1_HBGF
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Gencore version 4.5

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OM protein - protein search, using sw model

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May 22, 2002, 14:09:08; Search time 27.07 Seconds

(without alignments)
996.942 Million cell updates/sec

Title:

Perfect score:

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Sequence:

1 MAVLSGALCFRMKDSALKVL......IPEDPAWDAPITDFYFQQCD 156
Scoring table:
BLOSUM62
Gapop 10.0, Gapext 0.5
Searched:
562222 seqs, 172994929 residues
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otal number of hits satisfying chosen parameters: · 562222

Minimum DB seq length: 0 Maximum DB seq length: 2000000000 Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database : SPTREMBL_19:*

1: Sp_archea:*
2: Sp_bacteria:*
3: Sp_fungi:*
4: Sp_human:*
5: Sp_invercebrate:*
6: Sp_invercebrate:*
7: Sp_mncal:*
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9: Sp_phage:*
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073909 gallus gall Q9uha7 homo sapien Q9d&z6 mus musculu Q9nxH8 homo sapien O9nxEs xenonus lae	091wp7 mms musculu 029082 sus serofa 099043 oncorhynchu 091z15 sigmodon hi	Q9ddf3 cyprinus ca Q9ddf2 cyprinus ca Q9ttk1 turslops tr Q9ttk1 turslops tr Q9ptL2 oncorrhynchu Q96fb5 homo sapien	090w84 dicentrarch 057398 cyptinus ca 09pw18 cyptinus ca 09tsj0 equus cabal 09tsj0 equus cabal 09tsj8 scophthalmu	091562 xenopus lae 091562 xenopus lae 061155 rattus norv 095262 leishmanla 091679 dicryostell 093748 thermus the 095013 streptomyce 049364 arabidopsis 094962 rattus norv
169.5 20.7 267 13 167.5 20.5 158 4 165.5 20.2 183 11 157 19.2 169 4 143 17.5 289 4	128 15.6 599 11 127 15.5 267 6 Q 126 15.4 260 13 125 15.3 267 11	15.0 15.0 14.9 14.9	119 14.5 261 13 117 14.3 276 13 117 14.3 276 13 108.5 13.2 118 6 88.5 10.8 64 13 84 10.3 259 16	10 13

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Suzuki H., Toyo-oka K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
Whishaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
Hayashizaki Y.;
R. Hayashizaki Y.;
R. Functional annotation of a full-length mouse cDNA collection.";
R. Nature 409:685-690(2001).
R. EMBL; AK090941; BAB26471.1; -.
R. EMBL; AK090941; BAB2602.1; -.
R. EMBL; AK008975; Interleukin_1.
R. EMBL; AK008975; Interleukin_1.
R. EMBL; PEPPO; IPR000975; Interleukin_1.
R. EMBL; PEPPO; IPR000955; Interleukin_1.
R. EMBL; PEPPO; ISPS Interleukin_1.
R. EMBL; PEPPO; ISPS Interleukin_1.
R. EMBL; PEPPO; ISPS Interleukin_1.
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Barton J.L., Nicklin M.J.H.;

"IL-1Ll: A Novel Member of the Interleukin-1 Gene Family is Expressed in Trophoblasts and Macrophages.";

Submitted (OCT-1999) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             foung P.R.; Identification and initial characterization of four novel members
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G.I., Doyle M.L.,
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Mus.
NCBL_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 156;
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Kumar S., McDonnell P.C., Lehr R., Tierney L., Tzimas M.N.
Griswold D.E., Capper E.A., Tal-Singer R., Wells G.I., Doy
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                                                                                                                                                                                                                                                                                                                                                                                       000125; ILI; 1.
PS00253; INTERLEUKIN_1; 1.
156 AA; 17136 MW; A4D1EE2F93CF77A7 CRC64;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
IL-ILI PROTEIN (INTERLEUKIN-1 HOMOLOG 3).
ILIFS OR ILIHYI.
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100.0%; Pred. No. 5.2e-80;
ive 0; Mismatches 0;
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J. Biol. Chem. 275.10308-10314(2000).
EMBL; A7250429; CAB5983.1.1; -.
EMBL; AF200495; AAF69251.1; -.
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SEQUENCE
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Debets R., Timens J.C., Zurawski S., Sana T.R., Bazan F.,
Asstelein R.A.;
"Novel IL-1 ligands IL-1d and IL-1e use IL-1R related protein 2.";
Submitted (FEB-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                       2 MVLSGALCFRAKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSPVILG
                                                                                                                                                                           1 MYLSGALCFRWKDSALKYLYLHNNQLLAGGLHAEKVIKGEEISVYPNRALDASLSPVILG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
FILL DELTA (INTERLEUKIN-1 LIKE PROTEIN 1) (INTERLEUKIN-1 RECEPTOR
ANTAGONIST HOMOLOG 1) (INTERLEUKIN-1 DELTA).
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Mulero J.J., Nelken S.T., Ford J.E.;
"Organization of the Human Interleukin-1 Receptor Antagonist Gene
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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"A tissue specific interleukin-1 receptor antagonist homolog from
IL-1 cluster lacks IL-1, IL-1ra, IL-18 and IL-18ra activities.";
Submitted (JAN-2000) to the EMBL/GenBank/DDBJ databases.
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Barton J.L., di Giovine F.S., Symons J.A., Nicklin M.J.H.;
"A tissue specific interleukin-1 receptor antagonist homolog III cluster lacks IL-1, II-1ra, IL-18 and IL-18ra activities. Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        К.Е.,
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MEDLINE=99443727; PubMed=10512743;
Mulero J.J., Pace A.M., Nelken S.T., Loeb D.B., Correa T.R.,
Mulero J.S., Ford J.E.;
"ILLIHT: A Novel Interleukin-1 Receptor Antagonist Gene.";
Blochem. Biophys. Res. Commun. 263:702-706(1999).
   Length 155;
                                                                   Indels
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Smith D.E., Renshaw B.R., Ketchem R.R., Kubin M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Four New Members Expand the IL-1 Superfamily.";
   Score 814; DB 11;
Pred. No. 1.8e-79;
                                                                                                                                                                                                                                                                                                                                                                                                                122 LCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                       121 LCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           155 AA
99.4%; Score 814; DE
ilarity 100.0%; Pred. No. 1.8
Conservative 0; Mismatches
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SEQUENCE FROM N.A.
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   Query Match
Best Local Simil
Matches 155; (
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Receptor. SEQUENCE

DR NO DR NO

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Tursiops truncatus (Atlantic bottle-nosed dolphin).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Cetacea; Odontoceti; Delphinidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         68 CLSC---GTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCT 124
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"Molecular cloning and functional expression of bottle-nosed dolphin (Tursiops truncatus) interleukin-1 receptor antagonist.";
Vet. Immunol. Immunopathol. 78:131-141(2001).
HSSP; P18510; IIRA.
InterPro; IPR000975; Interleukin_1.
                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Strausberg R.; Strausberg R.; Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases. EMBL; BC009745; AAH09745.1; -.
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(TrEMBLrel. 19, Last sequence update)
(TrEMBLrel. 19, Last annotation update)
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(TrEMBLrel. 16, Last sequence update)
(TrEMBLrel. 19, Last annotation update)
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48.0%; Pred. No. 4.1e-25;
ive 16; Mismatches 45;
                 120 WFLCGPAEPQQPVQLTKESEPSA----RTKFYFEQ 150
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-131 AMEADQPVSLTNMPDE---GVMVTKFYFQE 157
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ProDom; PD002536; Interleukin_1; 1.
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Best Local Similarity 48.0%
Matches 72; Conservative
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SEQUENCE
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MEDLINE-21282553; Pubmed-11278614;
Lin H., Ho A.S., Haley-Vicente D., Zhang J., Bernal-Fussell J.,
Pace A.M., Hansen D., Schweighofer K., Mize N.K., Ford J.E.;
"Cloning and characterization of ILLHY2, a novel interleukin-1 family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61 VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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Pred. No. 6.8e-71;
4; Mismatches 10; Indels
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EMBL; AF334756; AAK68049.1; -.
EMBL; AF334755; AAK68048.1; -.
SEQUENCE 152 AA; 16975 MW; DF9735100FC711B2 CRC64;
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ProDom; PD002536; Interleukin_1; 1.
                                                                                                                                                                                                                  InterPro; IPR000975; Interleukin_1
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EMBL, AF186094, AAF25210.1; --
EMBL, AJ242731; CAB59822.1; --
EMBL, AJ242738; CAB59822.1; --
EMBL, AJ242738; CAB59823.1; --
EMBL, AJ271338; CAB5704.1; --
EMBL, AF230377; AAF91274.1; --
HSSP; P18510; 117.8
                                                                                                                                                                                                                                                                                                                                                                                 155 AA; 16962 MW;
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Best Local Similarity 91.0%;
Matches 141; Conservative
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tes 70; Conserv
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Q969H5; Q969H5

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RESULT Q969H5 ILLHY2

Query Match Best Local S

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Gaps

Indels 17; Length 159;

us-09-770-528-2.rspt

Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.

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Eukaryota, Metazoa, Chordata,
Mammalia, Eutheria, Primates,
                      Homo sapiens (Human).
                                                                                NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                               Receptor.
SEQUENCE
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SEQUENCE
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                                                                                                                                                                                                                                  68 CLSC---GTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCT 124
                                                                                                                                                                                                              CLSCGTEKGPILK--LEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTS 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                    10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSP--VILGVQGGSQ 67
                                                                                                                            10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSP--VILGVQGGSQ 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shin I.-S., Youn H.-Y.; "Molecular cloning of canine interleukin-1 receptor antagonist (IL-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalla; Eutheria; Carnivora; Fissipedia; Canidae; Canis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17;
                                                                                  15;
                                        Length 177;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           34.9%; Score 285.5; DB 6; Length 176; 46.0%; Pred. No. 1.1e-22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     48; Indels
                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lra).";
Submitted (JAN-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, AV026462; AAK01472.1;
HSSP, P18510; ILLR.
InterPro; IPR000975; Interleukin_1.
Pfam; PF00340; ILL; 1.
ProDom; PD002536; ILL; 1.
SNART; SM001253; ILL; 1.
SNART; SM0012536; ILL; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE 176 AA; 19922 MW; 8486CA54A240212B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q9BEH0;
01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OBBXX1 PRELIMINARY; PRT; 144 AA.
O9BXX1;
01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST FKSG75.
                                     37.3%; Score 305.5; DB 6;
47.7%; Pred. No. 7.7e-25;
tive 15; Mismatches 48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                 176
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                                                                                                                                                                                                                                                                                                                       150 LETDQPVGLTNTPQDA---VQVTKFYFQQ 175
                                                                                                                                                                                                                                                                                                 126 PEADQPVRLTQIPEDPAWDAPITDFYFQQ 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     69; Conservative
                                                                                Conservative
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SEQUENCE FROM N.A. Wang Y.-9, Li T., Gong L.; Wang Y.-9, Li T., Gong L.; "Identification and characterization of FKSG75, a novel member of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             68 CLSC---GTEKGPILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCT 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             41 EEISVVPNRALDASLSPVILGVQGGSQCLSC-GTEKGPILKLEPVNIMELYLGAKESKSF 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostoml;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae; Canis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17;
                                                                                                                                                                                                                                                                                                                                                                                     Length 144;
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Campbell S.E., Nasir L., Argyle D.J., Gault E.A., Duthie S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             49; Indels
                                                                          interleukin-1 family.";
Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases.
EMBL, AY005673; AAK01948.1;
HSSP; P18510; 1IRP.
Interpro; IRP00975; Interleukin_1.
Pfam: PF00340; Ill.1; 1.
Probom; PP0002536; Interleukin_1: 1.
                                                                                                                                                                                                                                                                                                           8733B16D6F118BB9 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  176 AA; 19938 MW; 8486CA54A254206B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST.
                                                                                                                                                                                                                                                                                                                                                                                   DB 4;
                                                                                                                                                                                                                                                                                                                                                                                   34.2%; Score 280.5; DB 4;
49.6%; Pred. No. 2.9e-22;
Live 19; Mismatches 34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
33.6%; Score 275.5; DB 6
Best Local Similarity 45.3%; Pred. No. 1.3e-21;
Matches 68; Conservative 16; Mismatches 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bennett D.;
"Cloning of canine IL-lra, INFR and TIMP-2.";
Vet. Immunol. Immunopathol. 78:207-214(2001).
HSSP; P18510; 1II.R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROSITE; PS00253; INTERLEUKIN_1; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pfam; PF00340; IL1; 1.
ProDom; PD002536; Interleukin_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IPR000975; Interleukin_1.
                                                                                                                                                                                                                                                                                                             144 AA; 15811 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                      57; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRELIMINARY;
                                                                                                                                                                                                                                                     SMART; SM00125; IL1; 1.
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Best Local Similarity
Matches 57; Conserv
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NCBI_TaxID=9606;
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SEQUENCE
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Q9UHA6
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64 GGSQCLSCGTEKG-PILKLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFL 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASLSP-----VILGVQ 63
                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Perissodactyla; Equidae; Equus.
NCBI_TaxID=9796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   11;
                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
MEDLINE-20092888: PubMed-10625660;
Smith D.E., Renshaw B.R., Ketchem R.R., Kubin M., Garka K.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                          24.2%; Score 198.5; DB 4; Length 157; 36.6%; Pred. No. 2e-13; Live 22; Mismatches 52; Indels 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
STRAIN-BREED THOROUGHBRED;
Dhar A.K., Thompson M.S., Paradis M.R., Alcivar-Warren A.;
"Molecular Characterization of Equine Interleukin 1 Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-NOV-1998 (TrEMBLrel. 08, Created)
01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST SECRETORY FORM (FRAGMENT)
                                                                                                                                                                                                                                                                                                                                                                          PROSITE; PS00253; INTERLEUKIN_1; UNKNOWN_1.
SEQUENCE 157 AA; 17702 MW; 7A54F3D7557A3EE3 CRC64;
                                                                                                                                                  Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                        Sims J.E.;
"Four New Members Expand the IL-1 Superfamily.";
J. Biol. Chem. 275:1169-1175(2000).
EMBL: AF201833; AAF25213.1;
HSSP; P10749; 2MIB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            72 AA.
                                                                                                                  157
                                                : |||: | || ||:
149 ALEADRLVSLINRPEEAMM---VIKFYFQK 175
                                 125 SPEADQPVRLTQIPEDPAWDAPITDFYFQQ 154
                                                                                                                                    01-MAY-2000 (TrEMBLrel. 13, Created) 01-MAY-2000 (TrEMBLrel. 13, Last seq 01-OCT-2001 (TrEMBLrel. 18, Last anno
                                                                                                                                                                                                                                                                                                                                         Pfam; PF00340; III; 1.
ProDom; PD002536; Interleukin_1; 1.
SMART; SM00125; IL1; 1.
                                                                                                                 PRT;
                                                                                                                                                                                                                                                                                                                                 InterPro; IPR000975; Interleukin_1.
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Best Local Similarity 36.6'
Matches 49; Conservative
                                                                                                               PRELIMINARY;
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                                                                                                                         Q9UHA5;
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                                                                                         RESULT 10
Q9UHA5
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077771
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82 EPVNIMELYLGAKESKSFTFYRDMGLTSSFESAAYPGWFLCTSPEADQPVRLTQIPEDP 141
                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                              Ouery Match 23.1%; Score 189.5; DB 4; Length 192; Best Local Similarity 37.3%; Pred. No. 2.4e-12; Matches 50; Conservative 23; Mismatches 50; Indels 11; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10 FRMKDSALKVLYLHNNQLLAGGLHAEKVIKGEEISVVPNRALDASL---SPVILGVQGGS 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                              Length 72;
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MEDLINE-20092888; Pubmed-10625660;
Smith D.E., Renshaw B.R., Ketchem R.R., Kubin M., Garka K.E.
                                                                                                                                                                                                                                                                                                                                          Indels
Pfam; PF00340; IL1; 1.
ProDom; PD002536; Interleukin_1; 1.
SMART; SM00125; IL1; 1.
SEQUENCE 192 AA; 21543 MW; 4AF584C81802F612 CRC64;
                                                                                                                                                                                                         1 1
72 AA; 8215 MW; 290CC9B9D4C413D9 CRC64;
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Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                         Query Match 23.3%; Score 190.5; DB 6; Best Local Similarity 53.4%; Pred. No. 5.4e-13; Matches 39; Conservative 11; Mismatches 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Four New Members Expand the IL-1 Superfamily."; J. Blol. Chem. 275:1169-1175(2000). EMB!, AF201832, AAF25212.1; -- HSSP; P18510; 1ILR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      192 AA
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                                                                                                                                                          PROSITE; PS00253; INTERLEUKIN_1; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
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151 ICTSCNCNEPVGVT 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           142 AWDAPITDFYFQQ 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 --SEMVTKFYFQE 70
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Homo sapiens (Human).
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5

Gaps

11;

Length 218;

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MEDLINE-21066552; PubMed-11145836; Pan G., Risser P., Mao W., Baldwin D.T., Zhong A.W., Filvaroff E., Pan G., Risser P., Mao W., Baldwin D.T., Zhong A.W., Vandlen R.; "Yansura D., Lewis L., Epenbrot C., Henzel W.J., Vandlen R.; "IL-1H, an interleukin l-related protein that binds IL-18 receptor/IL-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identification and initial characterization of four novel members of
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MEDILINE=21085660; PubMed=11217851;
Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,
Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
Aizawa K., Izawa M., Nishii K., Kiyosawa H., Kondo S., Yamanaka I.,
Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
MEDLINE=20209405; PubMed=10744718;
Kumar S., McDonnell P.C., Lehr R., Tierney L., Tzimas M.N.,
Griswold D.E., Capper E.A., Tal-Singer R., Wells G.I., Doyle M.L.,
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Debets R., Timans J., Zurawski S., Bazan J.F., Kastelein R.A.;
Novel IL-1 family member IL-1e responds through the orphan IL-1R-
related protein 2; response is antagonized by IL-1d.";
Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (Mouse).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-0cr-2000 (TrEMBLrel. 15, Created)
01-0cr-2000 (TrEMBLrel. 15, Last sequence update)
01-0cr-2001 (TrEMBLrel. 19, Last annotation update)
11TRELEUKIN-1 HOWOLGG 1 (INTERLEUKIN-1 EPSILON) (INTERLEUKIN 1 SUPERRAMILY 1, EPSILON).
11LIF6 OR FILL OR ILIE.
                                                                                                                                                                                                                                                                                                                                                                                                                        Ouery Match 23.1%; Score 189.5; DB 4; Length 2 Best Local Similarity 37.3%; Pred. No. 2.8e-12; Matches 50; Conservative 23; Mismatches 50; Indels
                                                                                                                                                                                                                                              Interpro; IPR000975; Interleukin_1.
Pfam; PF00340; IL1; 1.
Probom; PD002536; Interleukin_1; 1.
SEQUENCE 218 AA; 24138 WW; 76E09C35093DEA63 CRC64;
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J. Biol. Chem. 275:10308-10314(2000).
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                                                                                                                                                  1Rrp.";
Cytokine 13:1-7(2001).
EMBL; AF251119; AAG14421.1;
HSSP; P18510; 11LR.
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SEQUENCE FROM N.A.
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                                               01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel: 19, Last annotation update)
INTERLEUKIN-1 HOMOLOG 4 (IL-1X PROTEIN) (INTERLEUKIN-1-RELATED PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pan G., Risser P., Mao W., Baldwin D.T., Zhong A.W., Yansura D.,
Lewis L., Elgenbrot C., Henzle W.J., Vandlen R., Filvaroff E.,
"ILL-1H, an interleukin-1-related protein that binds IL-18 receptor/IL-
IRrp.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         60 FSIHDODHKVLVLDSGNLIA--VPDKNYIRPEIFFALASSLSSASAEKGSPILLGVSKGE 117
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                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

MEDISONE-FERTAL LUNG.

MEDLINE-20209405; Pubmed-10744718; FETAL B-CELL, AND FETAL COLON;

MEDLINE-20209405; Pubmed-10744718;

Kumar S., McDonnell P.C., Lehr R., Tierney L., Tzimas M.N.,

Griswold D.E., Capper E.A., Tal-Singer R., Wells G.I., Doyle M.L.,
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Wataryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23.1%; Score 189.5; DB 4; Length 218; 37.3%; Pred. No. 2.8e-12; Live 23; Mismatches 50; Indels 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF200496; AAF69252.1; -.
EMBL; AF167368; AAG29344.1; -.
EMBL; AF167118; AAG14420.1; -.
HSSP; P18510; 11LR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Manoj P.P., Mantovani A., Muzio M.;
Submitted (JUL-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 218 AA; 24126 MW; 96E089310D2CEA68 CRC64;
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Last annotation update)
218 AA.
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01-MAR-2001 (TrEMBLrel. 16, Last sequence v
01-DEC-2001 (TrEMBLrel. 19, Last annotation
INTERLEUKIN-1-RELATED PROTEIN LONG ISOFORM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the interleukin-1 family.";
J. Biol. Chem. 275:10308-10314(2000).
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ProDom; PD002536; Interleukin_1; 1.
  PRT;
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Matches 50; Conservative
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PRELIMINARY;
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177 ICTSCNCNEPVGVT 190
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TISSUE-COLON CARCINOMA;
                                                                                                                                                  LONG ISOFORM A).
Homo sapiens (Human)
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                                                                                                                                                                                                                                                 NCBI_TaxID=9606;
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RA Schriml L.M., Staubli F., Suzuki R., Tonita M., Wagner L., Washio T.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
Lyons P., Marchhonni L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Sasaki H., Toyoroka K., Wang K.H., Waltz C., Shibata Y., Storch K.-F.,
RA Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
RH Hayashizaki Y.;
Runctional annotation of a full-length mouse cDNA collection.";
RE WEL; ARO04061; BAB23147.11; -.
DR EMBL; RO0340; ILL1 1.
PRODOM; PD002556; Interleukin_1; 1.

PRODOM; PD002556; Interleukin_1; 1.

SEQUENCE 160 AA; 18015 MW; AA0434D68FF62F4A CRC64;
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Search completed: May 22, 2002, 14:12:31 Job time: 203 sec

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